					Table 1-3					
53022		1.34	0.70	0.24	1.90	c ca 1				
377246	AA055306	12.60	3.02	3.78	4.17	5.63 3.34	-	-		
814662	AA481045	1.12	0.67	0.19	1.67	5.84	+	+	+	•
45327		8.18	2.30	2.08	3.56	3.94	•	•	+	+
51700	H22853	3.66	1.21	0.82	3.02	4.47	•	+	+	+
80050		8.15	3.84	1.52	2.12	5.35	-	+	+	+
594994	AA 172372	14.61	10.98	2.38	1.33	6.14	•	•	+	+
795907	AA460346	5.39	1.25	1.71	4.31	3.15	+	-	+	+
824126	AA490611	0.59	0.37	0.10	1.58	5.87		+	+	+
511952	AA100674	225.66	92.92	45.17	2.43	5.00	<del>-</del>	•	+	+
823615	AA496957	9.40	3.54	1.97	2.65	4.76	+	-	+	+
	AA432075	17.23	6.93	3.50	2.49	4.93	<del></del>	+	+	+
	AA918370	401.67	285.67	66.91	1.41	6.00	-	<b>-</b> ,	<b>+</b> .	+
1605407	AA988746	9.00	4.61	1.65	1.95	5.45	-	•	<b>+</b>	+
1606837	AA996028	23.76	9.09	4.98	2.61	4.77	•	-	+	+
768395	AA495819	14.29	3.54	4.28	4.03	3.34	- +	+	+	+
796239	AA460675	26.07	23.22	4.18	1.12	6.24	<i>T</i> •	+	<del>-</del>	+
49302		4.88	0.86	2.90	5.68	1.68	+	•	+	+
754358	AA436142	12.77	7.46	2.26	1.71	5.64	•	+	<del>-</del>	-
207029	R98842	81.88	14.79	45.41	5.54	1.80	+	-	+	+
838285	AA458747	1.34	1.36	0.21	0.99	6.35	-	+	-	-
1636108		3.58	1.45	0.74	2.47	4.85		•	+	+
	AA460719	0.66	0.83	0.10	0.80	6.50	-	-	+	+
238821	H65029	6.54	2.22	1.50	2.94	4.35	+	+	+	+
129024		4.30	1.04	1.36	4.13	3.16	+		+	+
	AA430718	16.14	6.63	3.34	2.43	4.83	-	+	+	+
289071	N63609	37.48	15.32	7.78	2.45	4.82	+	-	+	+
1637282		4.01	2.58	<b>0.</b> 70	1.55	5.71	<u>.</u>	-	+	+
263697		4.23	0.93	1.58	4.57	2.68	+	. <b>-</b> +	<del>.</del>	+
	AA683050	35.22	12.06	8.15	2.92	4.32	+	+	- +	+
32381		2.18	1.18	0.41	1.85	5.38	_	+	+	+
	AA130677	1.92	1.75	0.31	1.10	6.12	-	_	+	+
	AA465148	1.62	0.58	0.37	2.81	4.40	_	+	+	+
	AA074258	12.04	3.46	3.22	3.48	3.73	+	+	+	+
416374		4.49	2.66	0.81	1.69	5.52	-	-	+	+
	AA448402	12.06	3.84	2.97	3.14	4.05	+	+	+	+
213682		13.07	4.10	3.27	3.19	3.99	+	+	+	+
	AA884713	3.68	0.99	1.07	3.72	3.44	-	+	_	+
	AA032221	31.04	12.66	6.60	2.45	4.70	-	•	<b>.</b>	+
	AA055880	132.22	56.58	<b>27.</b> 52	2.34	4.81	-	-	+	+
242952		16.94	4.39	5.16	3.86	3.28	-	+	-	+
	AA775422	3.53	0.87	1.16	4.07	3.05	+	+	-	+
853968 287728		12.10	2.86	4.21	4.23	2.87	+	+	+	+
	N75893 AA486445	2.76	0.70	0.89	3.97	3.10	+	+	+	+
	AA676765	4.24	3.59	0.72	1.18	5.88	•	-	+	+
	AA670286	4.95	1.36	1.45	3.63	3.42	+	+	+	+
	AA011176	6.80	3.30	1.37	2.06	4.97	•	-	+	+
2 <b>727</b> 50	N36233	5.21	2.32	1.09	2.25	4.78	-	•	+	+
	AA490390	20.57	6.04	5.70	3.40	3.61	+	+	+	+
	AA598531	11.96	3.51	3.34	3.41	3.58	-	+	•	+
	AA663983	5.08	2.77	0.98	1.83	5.16	•	-	+	+
264895		50.50	12.03	18.24	4.20	2.77	+	+	-	-
	AA504120	7.86	1.63	3.70	4.82	2.12	+	+	-	-
JEJ201	JULY 204 150	11.96	2.92	4.19	4.09	2.86	+	+	•	-

Page 4

					Table 1-3					
461476	AA705041	0.78	0.61	0.14	1.28	5.64	-	-	+	+
323474	W45623	1.25	0.32	0.41	3.87	3.06	+	+	+	+
	AA625675	2.69	0.70	0.87	3.83	3.10	+	+	•	+
	AA401111	4.93	1.40	1.45	3.51	3.40	+	+	+	+
	AA709044	12.93	4.03	3.49	3.21	3.70	•	+	+	+
	AA777705	3.81	0.83	1.64	4.58	2.32	+	+	+	
878544	AA775863	19.20	6.78	4.73	2.83	4.06	+	+	+	+
755239	AA422139	17.93	9.29	3.62	1.93	4.95	. <del>-</del>	<u>.</u>	+	+
259973	N47829	1.66	1.95	0.28	0.85	5.99	-	-	+	· +
290370	N62301	0.62	0.48	0.11	1.30	5.53	-	-	+	+
773287	AA425534	15.94	8.17	3.27	1.95	4.87	•	-	+	+
	AA406505	7.78	2.14	2.45	3.63	3.18	_	+	•	+
290841	N99711	6.35	2.89	1.38	2.20	4.61	-	+	+	+
<b>5</b> 060 <b>6</b>	H17763	14.94	6.00	3.47	2.49	4.31	-	•	+	+
1472689	AA873159	5.15	1.21	2.03	4.26	2.53	+	+	+	
252314	H87175	0.69	0.62	0.12	1.11	5.66	-	•	+	+
813275	AA455940	1.75	0.86	0.37	2.04	4.73	-		+	+
951125	AA620556	18.86	7.00	4.63	2.69	4.07	+	•	+	+
454317	AA677165	20.56	33.95	3.34	0.61	6.16	-	•	+	+
433253	AA699427	7.29	2.43	1.95	3.00	3.74	+	+	+	+
1585549	AA976714	6.44	3.55	1.31	1.81	4.92	-	-	+	+
491764	AA150502	0.81	0.59	0.15	1.36	5.37	•	•	+	+
	AA133044	559.22	179.15	155.61	3.12	3.59	+	+	+	+
245547		5.86	1.63	1.88	3.59	3.12	+	+	4	+
200599	H48472	13.42	5.61	3.11	2.39	4.32	-	+	+	+
139771	R62352	7.72	6.17	1.42	1.25	5.44	-	-	+	+
	AA705376	8.32	6.33	1.55	1.31	5.38	-	-	+	+
	AA464580	4.37	1.58	1.11	2.76	3.92	-	+	+	+
279482	1	8.66	3.25	2.16	2.67	4.00	+	•	+	+
127396		14.07	5.12	3.61	<b>2.7</b> 5	3.90	~	+	+	+
	AA490609	4.64	2.75	0.94	1.69	4.95	-	•	+	+
	AA609323	38.01	9.06	15.59	4.20	2.44	+	+	+	-
	AA287375	9.94	3.00	3.00	3.31	3.31	-	+	-	+
2508 <b>69</b>	•	123.92	32.54	44.10	3.81	2.81	4	•	+	-
1631747		5.09	1.33	1.84	3.84	2.77	+	+	-	+
	AA779391	18.03	7.40	4.35	2.44	4.14	-	+	+	+
	AA609368	7.90	4.94	1.59	1.60	4.98	•	- ' '	+	+
81578		12.77	4.82	3. <b>25</b>	2.65	3.93	-	+	+	+
	AA434360	103.13	30.63	32.18	3.37	3.20	+	+	+	+
	AA621761	14.86	3.64	<b>5.98</b>	4.08	2.48	+	+	-	-
	AA459100	18.28	10.68	3.77	1.71	4.85	-	•	+	+
77915		14.99	17.48	2.63	0.86	5.70	-	-	+	+
260116		0.88	0.86	0.16	1.03	5.52	-	-	+	+
	AA458551	17.84	5.98	5.02	2.98	3.55	-	+	-	+
210405	1	34.02	7.84	15.49	4.34	2.20	+	+	-	-
124824	1	30.85	11.28	8.13	2.73	3.79	+	•	+	+
	AA169379	17.74	7.51	4.26	2.36	4.17	+	+	+	+
	AA278473	4.77	2.81	0.99	1.70	4.82	-	•	+	+
	AA058818	15.01	6.81	3.48	2.20	4.32	-	-	+	+
. 191603		22.14	4.64	12.72	4.77	1.74	+	+	-	-
	AA012866	40.29	13.93	11.17	2.89	3.61		+	+	+
	AA491265	0.74	0.41	0.16	1.81	4.68	-	•	+	+
	AA181179 AA885642	5.74	2.03	1.58	2.82	3.64	+	+	+	+
1200000	JAN 003042	6.11	3.34	1.32	1.83	4.63	-	•	+	+

							<del></del>			
								الأيدية الإسبال الأمر الإ	Hadi salla se' S	الكنيلا بطابت الدينيز بيسطة البين
•				Ti	able 1-3					
813973 AA45	5689 3	3.24 0	.88.	.18		2.75	-	+	•	•
825297 AA50						4.97	-	-	+	+
592403 AA15	9497	0.46			2.95	3.50	+	-	+	+
838807 AA45	7681				0.74	5.71	-	-	+	+
845352 AA64	4080 1				2.48	3.97	-	•	+	+
814798 AA46				1.58	2.10	4.34	-	-	+	+
1607765 AA98				0.96	2.29	4.14	+	+	+	+
1035765 AA62	29117			1.61	2.57	3.86	-	-	+	+
824739 AA48				83.0	2.57	3.86	-	+		+
	99659			2.28	3.02	3.40	_	+	•	+
				1.09	3.29	4.82	-	•	+	+
823691 AA4				1.54	1.59 2.82	3.58	•	+	-	+
	72336	2.10		0.59	1.03	5.36	-	-	+	+
1456160 AA8			10.32	7.76 0.98	4.25	2.12	+	+	-	•
897767 AAS		2.07	0.49	0.9 <b>8</b> 36.21	4.50	1.88	+	+	-	-
877641 AA4			<b>15</b> .10 3 <b>4.7</b> 3	3.28	2.60	3.76	+	-	+	+
504544 AA1		12.31 1.21	0.71	0.26	1.71	4.63	-	-	+	+
1466631 AA8		6.77	3.15	1.62	2.15	4.19	-	-	+	+
898109 AAS		0.53	0.50	0.10	1.05	5.28	-	-	+	+
280763 N 1492238 AA	150563 875036	13.96	4.83	4.08	2.89	3.43	-	+	-	+
	450770	5.44	1.19	3.10	4.56	1.76	+	+	-	-
462468 AA		3.24	0.96	1.10	3.36	2.96	-	+	-	+
768643 AA		41.57	23.92	9.10	1.74	4.57	-		+	+
	R40970	3.34	1.66	0.78	2.01	4.29	-	. •	+	+
610124 AA		1.70	1.73	0.32	0.98	5.32	-	-	+	+
	R39862	4.62	2.68	1.01	1.73	4.56	-	-	+	+
487499 AA		4.41	1.79	1.16	2.47	3.82	+	-	+	+
	1004315	25.01	12.24	5.91	2.04	4.24	-	-		<u>'</u>
204111	H55907	3.38	0.87	1.40	3.86	2.42	+	+	+	•
781454 A	4432312	9.75	3.24	2.99	3.01	3.26	+	+	+	+
491184 A		6.35	1.83	2.27	3.47	2.80	+	+	•	-
840788 A		23.03	5.46	11.26	4.22	2.05 3.86		· -	-	+
547058 A.		9.57	4.00	2.48	2.39	2.85		+	-	+
784253 A.		11.76	3.47	4.12	3.39 2.32	3.91		-	-	+
196189	R91950	11.19	4.82	2.86 4.18	4.30	1.93	+	+	-	-
47542	H16454	8.07	1.88	1.61	3.61	2.62		+	-	•
47384	H10778	4.22	1.17 31.45	19.54	2.38	3.83		-	+	+
824237 A		74.89 5.22	31.45	1.07	1.35	4.86		-	+	+
194723	R89849 H01205	5.29	2.31	1.35	2.29	3.92		•	+	+
150314	A486200	326.54	118.59	94.88	2.75	3.44	+	+	+	+
48286	H12190	2.34	0.61	0.99	3.83	2.36	;	+	•	-
	AA677287	2.69	0.70	1.13	3.82	2.38	3 +	+	-	-
24208		7.49	3.51	1.85	2.13	4.05		-	+	+
273546	N33274	9.10	3.09	2.81	2.95	3.23		+	-	+
267419	N24973	0.97	0.96	0.19	1.01	5.10		-	+	+ +
	AA412250		0.58	0.10	0.92			-	+	+
	AA977296		1.57	1.81	3.30			+	-	+
	AA505063	2.26	0.96	0.60				•	+	+
262864	l.		16.43	9.52				•	+	
	AA626146	30.25	19.90	6.55			1	•	+	
	A A 680244	51.58	35.87					. •	+	
3798 <b>0</b>	R61372	0.84	0.29	0.27	2.90	<b>3.</b> !	18   4	_		
										Page

Page 6

			<b>)</b>		Table 4 6		_			
124252	R02439	606	4.54		Table 1-3					
238661		6. <b>06</b> 2. <b>74</b>	4.74	1.25	1.28	4.85	•		+	+
	AA160080	1.08	2.18 1.02	0.56	1.26	4.86	-	•	+	+
785616	AA449004	11.47	3.12	0.21	1.07	5.04	•	-	<b>+</b> .	+
84022		6.97	2.71	4.70	3.67	2.44	-	+	-	-
	AA126951	7.85	2.71	1.97	2.57	3.54	-	•	+	+ '
34134	R44734	2.73	1.33	2. <b>46</b> 0.6 <b>8</b>	2.89	3.20	•	+	-	+
39885	R52543	0.94	0.30	0.8	2.06	4.03	-	-	+	+
840590	AA488014	21.70	7.54	6.7 <b>8</b>	3.11	2.98	-	+	-	+
49710	H28973	3.21	1.47	0. <b>78</b>	2.88	3.20	+	•	+	•
42803	R60014	2.96	18.0	1.23	2.18 3.66	3.89	•	-	-	+
134969	R31789	6.81	4.59	1.48		2.41	+	+	+	-
75415	T57609	51.71	24.46	13.09	1.48 2.11	4.58	-	-	+	+
843094	AA488626	49.18	13.00	21.59	3.78	3.95	•	•	-	+
137189	R36409	5.26	1.96	1.56	2.68	2.28	-	+	-	•
· 45578	H08016	5.05	1.63	1.72	3.10	3.37	•	+	•	+
430465	AA680186	5.55	1.53	2.30	3.62	2.94	-	+	-	+
67654	T49539	15.79	6.24	4.50	2.53	2.41	+	-	+	-
550141	AA102591	78.20	22.40	30 <b>.93</b>	3.49	3.51 2.53	-	-	+	+
796351	AA459765	2.34	1.80	0.50	1.30	4.72	+	+	-	•
1637296	AI005519	25.52	13.27	6.23	1.92	4.09	•	•	+	+
549101	AA083485	12.81	2.43	17.02	5.26	0.75	+	• ,	+	+
824376	AA489696	0.49	0.34	0.11	1.42	4.58	· ·	+	-	-
773204	AA425692	21.34	15.28	4.63	1.40	4.61	•	•	+	+
453599	AA679569	4.23	2.23	1.03	1.89	4.10	-	•	+	+
	AA481256	9.59	4.44	2.50	2.16	3.84	-	-	+	+
196650	R93237	10.93	6.59	2.52	1.66	4.33	_	•	+	+
725489	AA398521	26.71	9.34	8.54	2.86	3.13	_	+	+	+
1475987	AA872436	2.96	0.83	1.23	3.58	2.40	+	+	•	+
811162	AA485748	6.36	4.30	1.41	1.48	4.50		-	. <del>-</del>	-
300132	AA062805	6.22	1.68	2.75	3.72	2.26	-	+	-	+
701677	AA287067	16.03	15.09	3.27	1.06	4.91	<u>.</u> .		+	+
	AA442301	12.77	6.52	3.19	1.96	4.01	-	•	-	+
	AA788970	7.60	2.71	2.40	2.80	3.17	+	+	+	+
210862	H65659	2.56	1.16	0.68	2.21	3.76	-	•	+	+
777740	AA083671 AA425773	8.46	7.83	1.73	1.08	4.88	-	-	+	+
772416	AA405597	21.90	19.94	4.51	1.10	4.86	-	•	+	+
706512	AA460251	1.03	1.35	0.20	0.76	5.19	-		+	+
66902	T67444	17.83	7.09	5.20	2.51	3.43	-	+	+	+
	AA505162	3.21	1.14	1.03	2.82	3.12	+	+	+	+
201282	R99503	4.03	1.32	1.40	3.06	2.88	-	+		+
	AA987644	6.42	4.16	1.46	1.55	4.39	-	•	+	+
362926	AA018979	1.97 4.39	0.81	0.57	2.45	3.48	+	•	+	+
1456721	AA864875	12.82	2.04	1.16	2.15	3.77	-	-	+	+
1070015	AA599717	3.26	5.49	3.58	2.33	3.58	•	•	•	+
109153	T81140	5.46	1.43 1.70	0.90	2.28	3.63	•	+	+	+
50764		0.43		2.03	3.20	2.69	-	+	-	-
356992		21.95	0.23 7.93	0.11	1.89	4.00	-	•	+	+
	AA996104	6.06	7.93 2.17	7.04	2.77	3.12	-	+	-	+
824843	AA488865	4.03	2.24	1.97	2.79	3.07	+	•	+	+
259462	N29545	3.69	1.65	0.99	1.80	4.06	. •	•	+	+
	AA425205	26.40	17.78	1.02 6.05	2.24	3.62	+	•	+	-
626990	AA190850	7.39	3.82	6.05 1.89	1.49	4.37	-	-	+	+
•	,		J.UZ	1.07	1.93	3.91	-	•	+	+

						#*. y	السب الدراد اد	صد وينابرو فاوبوؤاد	البينا سلليج البندر سرطة الهراة
			Tab	le 1-3		" <b>T</b>			
743 <b>749\AA634291</b>	5.10	1.65 1.5	36 3.	10 2	.74	-	+	•	•
79000 T62164	6.24	3.53			.05	•	•	+	+
23576 T77385	3.58	1.59 1.6	00 2.		.57	-	-	<u>-</u>	+
366815 AA029490	71.61	18.49 36.			1	+	•	+	• +
731016 AA421256	0.85				1.96	-	-	+	•
415437 W81124	3.5 <b>2</b>	•	-		2.35	•	+	+	+
197779 R93515	8.61	_			4.50 4.25			+	+
1519515 AA902831	0.85				4.79	<u>.</u>	•	+	+
511428 AA126009	24.31				3.32	•	•	-	+
624785 AA181995	6.17	_			3.87	-	-	4.	+
809578 AA456616	24.75				4.01	-	-	+	+
360403 AA015658	28.16			3.01	2.77	•	+	-	+
431286 AA682627	13.81 0.50			1.68	4.10	•	-	+	+
712216 AA280288				4.85	0.93	+	+	-	•
296788 W01113 1638852 A1016618				2.41	3.36	•	-	•	+
1636868 A1015589				1.76	4.01	-	-	+	+
461592 AA705142				0.92	4.84	-	-	+	+ ·
357364 W9370			0.23	2.67	3.09	-	•	+	+
594517 AA16964		12.50	6.57	1.98	3.78	-	-	+	+
842882 AA48641		1.38	0.53	1.60	4.16	-	-	+	+
111812 T8497		1.89	1.98	2.94	2.81	-	+	+	· +
757206 AA44396	<b>9</b> 5. <b>25</b>	4.09	1.17	1.28	4.47	-	+	-	-
292042 W0233		2.67	4.19	3.51	2.24	- -	+	+	+
743828 AA63437		1.15	1.17	2.90	2.85	+	-	· <u>-</u>	+
884657 AA6299	6.68	2.72	2.04	2.46	3.28	-	•	-	+
743804 AA6343		1.90	1.66	2.68 2.44	3.30	-	_	+	+
362829 AA0194		1.24	0.92 1.39	2.80	2.93	-	+	•	+
502690 AA1271		1.46 2.55	1.89	2.44	3.29	+	-	+	-
854678 AA6300		2.33 8.27	4.20	1.93	3.79	-	-	4	+ .
626385 AA1891		1.53	1.39	2.72	3.00	-	-	+	+
364022 AA0215 264465 N205		0.84	0.16	0.91	4.79	-	-	+	+
232628 H725			1.58	3.16	2.55	+	+	•	+
950410 AA599			1.49	3.56	2.14	+	~		- +
838761 AA457			0.58	1.23	4.47	•	-	+	+
753162 AA400		1.76	0.69	1.60	4.10	-	+	- `	+
610011 AA169	192 34.63		13.23	3.07	2.62	•	+	_	+
251330 H97	880 9.21		2.83	2.44	3.25 2.85	1 :	+		+
823588 AA497	127 7.72		2.70	2.83	3.83	1		+	+
361048 AA017			0.82	1.85 2.78	2.89	+	•	+	-
826102 AA52	4.1		1.45 18.48	3.25	2.40		+	•	•
877832 AA62	5 <b>628</b> 44.4		6.13	4.22	1.44		+	-	•
856447 AA63			0.71	3.73	1.93		+	-	<del>-</del>
	-		0.51	1.17	4.49		-	+	+
815549 AA45 768443 AA49			6.79	1.73	3.92	-	-	+	+
1456602 AA86			1.88	2.40	3.25		-	-	
745103 AA62			2.98	2.67	2.97		+	-	
784258 AA4			5.04	3.64	2.00		+	-	
950836 AA6		-	11.31	2.56			•	7 4	
609052 AAI	<b>76607</b> 0.	87 0.87	0.19	0.99			•		+ +
1637343 A10	1 <b>5359</b> 5.	.03 4.49	1.12	1.12			-		+ <b>-</b>
462325 AA7	05516 8	.76 3.83	2.63	2.29	3.3	٠ ١ ٠	_		
•									Page 8

Page 8

					Table 1-	3				
878284	AA670280	4.41	2.26	1.20.	1.95	3.66	ı			
486538	AA042812	7.64	5.06	1.86	1.51	4.10		-	+	+
	AA670296	5.34	1.86	1.95	2.87	2.73	•	•	+	+
251019		45.62	22.03	12.95	2.07	3.52	•	+	•	+
258167		17.24	8.15	4.96	2.12	3.48	•	-	+	+
416833		6.07	2.16	2.18	2.81	2.78	•	-	+	+
186768		9.43	5.75	2.39	1.64	3.95	-	+	-	+
1558411	AA976063	8.20	5.56	1.99	1.47	4.11	-	-	+	•
49728		1.27	0.57	0.38	2.22	3.36	•	-	+	+
125722		3.28	1.21	1.14	2.72	2.87	•	-	-	+
613237	AA181767	1.07	0.78	0.25	1.37		-	+	-	+
	AA112515	1.14	0.40	0.41	2.82	4.22	•	•	+	+
79032		37.58	31.85	8.53	1.18	2.77	•	+ .	-	+
843139	AA485922	39.96	18.13	11.84	2.20	4.40 3.37	•	-	. +	+
841302	AA487213	7.41	3.64	2.09	2.04		-	-	-	+
191978		47.95	29.72	12.11	1.61	3.54	+	-	+	•
135900		9.44	5.00	2.57	1.89	3.96	-	-	+	+
713109	AA282983	2.44	0.98	0.79	2.48	3.68	-	-	+	+
626640	AA191512	8.33	4.14	2.35	2.01	3.08	•	-	•	+
322914	W45148	9.34	3.68	3.10	2.54	3.54	•	-	+	+
251517	H96605	0.73	0.80	0.16	0.91	3.02	-	-	-	+
840878	AA482228	25.81	27.85	5.59	0.93	4.64	•	-	+	+
626716	AA191245	21.60	13,98	5.41	1.55	4.61	•	-	+	+
291345	W02978	70.40	31.69	21.23	2.22	3.99	-	-	+	+
80500	T64625	140.81	52.48	49.33	2.68	3.32	+	-	+	+
815737	AA485201	3.64	2.20	0.94	1.65	2.85	-	+	+	+
436769	AA703079	168.72	131.76	39.68	1.28	3.88	-	-	+	+
461509	AA705077	0.62	0.38	0.16	1.66	4.25	-	-	+	+
	AA001234	4.05	3.19	0.95	1.27	3.87	-	-	+	+
137638	R37224	5.06	1.71	1.97	2.95	4.25	-	-	+	· +
855624	AA664101	17.75	8.17	5.31	2.17	2.57	-	+	-	+
109316	T80924	12.35	4.01	5.08	3.08	3.34	+	· -	+	•
725978	AA399198	4.92	2.96	1.28	1.66	2.43	+	+	+	-
392678	AA708342	34.95	10.61	15.79	3.29	3.85 2.21	•	-	+	+
120707	T95670	19.60	12.65	4.95	1.55	3.96	-	+	-	-
841220	AA486723	1.46	1.23	0.34	1.18	4.32	•	-	+	+
1631194	AA994757	21.83	14.39	5.47	1.52	3.99	-	-	-	+
767798	AA418755	5.70	1.98	2.18	2.88	2.61	-	•	+	+
251435	H97989	0.73	0.90	0.16	0.81	4.68	~	+	•	+
75044	T51904	3.46	0.93	1.96	3.72	1.77	-	-	+	+
743071	AA405984	1.89	1.02	0.52	1.85		-	+	-	-
490766	AA133165	3.48	1.03	1.66	3.38	2.10	-	-	+	+
344589	W73144 .	53.28	21.92	17.55	2.43	3.04	•	+	•	•
250673	H959 <b>78</b>	2.66	1.12	0.86	2.37	3.09	•	+	-	+
785910	AA449644	4.01	1.01	2.72	3.98	1.48	+	•	+	•
824329	AA489670	1.94	1.47	0.47	1.32	4.13	+	+	-	- '
1420790	AA826237	13.40	6.84	3.84	1.96		<del>-</del>	•	+	+
430318	AA010608	10.87	4.02	3.96	2.71	3.49 2.74	-	•	•	+
45564	H05264	1.66	0.76	0.51	2.17	3.28	•	+	-	+
251936	H97488	7.48	2.74	2.76	2.73	2.71	-	-	+	+
253314	H89330	1.52	1.34	0.35	1.14	4.30	•	+	•	•
490232	AA121428	2.68	0.85	1.17	3.15	2.28	-	-	+	+
855244	AA630545	15.78	6.78	5.09	2.33	3.10	-	+	•	•
898242	AA598621	10.34	7.74	2.53	1.34	4.09	+	+	+	+
					٠.٠٠	4.09	-	•	+	+

195602  A757604   IA.45						Table 1-3					
264640			14.45	6.11		2.37		. •			
1031598   AA609474   5.18   1.34   3.37   3.88   1.54   +		AA417911	0.44	0.45		0.98		-	-	+	+
785913   AA4496577   3.35   0.80   2.75   4.19   1.22   + +   141505   R72977   6.38   1.86   3.21   3.42   1.99   - +   141505   R72977   6.38   1.86   3.21   3.42   1.99   - +   141505   R72977   6.38   1.86   3.21   3.42   1.99   - +   1475283   AA625567   7.94   5.76   1.97   1.38   4.02   +   1475283   AA625567   7.94   5.76   1.97   1.38   4.02   +   1473851   AA629801   7.71   4.73   2.05   1.63   3.76   + +   148531   AA629801   7.71   4.73   2.05   1.63   3.76   + +   148531   AA629801   7.71   4.73   2.05   1.63   3.76   + +   1495032   W76319   20.78   9.11   6.69   2.28   3.11   + +   1495034   AA65228   14.81   9.71   3.84   1.52   3.85   + +   151917   AA465228   14.81   9.71   3.84   1.52   3.85   + +   151917   AA465228   14.81   9.71   3.84   1.52   3.85   + +   1473300   AA916323   12.44   4.26   5.09   2.92   2.44   + +     1473300   AA916323   12.44   4.26   5.09   2.92   2.44   + +     1473300   AA916323   12.44   4.26   5.09   2.92   2.44   + +     147518   AA702247   10.98   6.11   3.08   1.80   3.56   +   +   1475190   AA42604   5.86   3.56   1.58   1.65   3.71   +   +   1475190   AA46643   6.44   3.83   1.75   1.68   3.68   +   +   1475190   AA765272   2.98   2.09   0.53   1.08   4.27   - +   +   +   1475190   AA765072   2.26   2.09   0.53   1.08   4.27   - +   +   +   121436   T97457   5.37   6.46   1.46   0.99   4.35     +   +   +   121436   T97457   5.37   6.46   1.46   0.99   4.35     +   +   +   121436   T97457   5.37   6.68   0.87   1.00   0.47   1.65   3.69     +   +   +   121436   T97457   5.37   6.68   1.69   0.91   0.				6.10				-	-	+	+
343987   W70234   11.55   12.13   2.59   0.95   4.46   -								+	+	-	•
141505		AA449657						+	+	-	-
298155   N70794   2.85   1.29   0.89   2.21   3.19   -   -   +   +   +   +   +   +   +   +						0.95		•	-	+	+
745283   AA625367		R72977						•	+	•	-
26230								-	-	-	+
743851         AA634424         8.22         3.22         2.89         2.55         2.84         -         +         -         +         848431         AA629801         7.71         4.73         2.05         1.63         3.76         -         -         +         +         +         72993         AA412049         4.95         3.66         1.23         1.35         4.03         -         -         +         +         +         729933         AA412049         4.95         3.66         1.23         1.35         4.03         -         -         +         +         +         73345         AA62528         1.81         9.71         3.84         1.52         3.85         -         -         -         +         +         773345         AA423442         1.21         1.11         0.28         1.09         4.29         -         -         +<								-	-	•	
88451   AAC29801   7.71   4.73   2.05   1.63   3.76   -								-	-	•	
345022 W76319 20.78 9.11 6.69 2.28 3.11 + + + + + + + + + + + + + + + + +								-	+	-	
729953 AA412049								-	<del>-</del> .	+	
815017   AA465258   14.81   9.71   3.84   1.52   3.85   + +   +								-	•	-	
308466   N95495   9.26								-	-	+	
T73345		2						-	-	-	
811766		1						•	-		
246869		1						•			
1473300       AA916323       12.44       4.26       5.09       2.92       2.44       +       +       -       -       -       47818       AA702247       10.98       6.11       3.08       1.80       3.56       -       -       +       +       +       -       -       +       +       +       293243       N68686       13.17       5.84       4.25       2.26       3.10       -       +		1						-		+	+
447518   AA702247         10.98         6.11         3.08         1.80         3.56         -         -         +         +         784093   AA446643         6.44         3.83         1.75         1.68         3.68         -         -         -         +         +         +         293243         N68686         13.17         5.84         4.25         2.26         3.10         -         +										-	-
784093 AA446643 6.44 3.83 1.75 1.68 3.68 + + 293243 N68686 13.17 5.84 4.25 2.26 3.10 - + + + + + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 - + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 + + 755409 AA424706 7.81 2.43 3.67 3.22 2.13 + + 755409 AA439121 1.74 1.05 0.47 1.65 3.69 + + + - + 755409 AA439121 1.74 1.05 0.47 1.65 3.69 + - + 755409 AA439121 1.74 1.05 0.47 1.65 3.69 + + 755409 AA599078 7.48 3.82 2.21 1.96 3.38 + + + 755409 AA599078 7.48 3.82 2.21 1.96 3.38 + + + 755409 AA599078 7.48 3.82 2.21 1.96 3.38 + + + 755409 AA599078 7.48 3.82 2.21 1.96 3.38 + + + 755409 AA599078 7.48 3.20 1.10 1.37 3.97 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.57 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA6462631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + + 755409 AA682631 1.85 1.07 0.51 1.73 3.59 + + + + + + + + + + + + + + + + +								+	+	-	
293243 N68686 13.17 5.84 4.25 2.26 3.10 - + + + +   897733 AA598996 5.86 3.56 1.58 1.65 3.71 + +   755409 AA424706 7.81 2.43 3.67 3.22 2.13 - +   299723 N75055 2.98 2.09 0.76 1.43 3.93 + +   461499 AA705072 2.26 2.09 0.53 1.08 4.27 + +   82905 T69305 1.21 0.48 0.43 2.50 2.85 - + - +   824917 AA489121 1.74 1.05 0.47 1.65 3.69 +   121436 T97457 6.37 6.46 1.46 0.99 4.35 + +   809657 AA5999078 7.48 3.82 2.21 1.96 3.38 +   809657 AA456331 4.72 2.68 1.32 1.76 3.57 +   145388 R78386 2.66 0.94 1.06 2.83 2.50 - + -    145388 R78386 2.66 0.94 1.06 2.83 2.50 - + -    145388 R78386 2.66 0.94 1.06 2.83 2.50 - + -    145388 R78386 2.66 0.94 1.06 2.83 2.50     145388 R78386 2.66 0.94 1.06 2.83 2.50     145388 R78386 2.66 0.94 1.06 2.83 2.50		•						-	-	•	
897733 AA598966		•					1	•			
755409 AA424706 7.81 2.43 3.67 3.22 2.13 - +		2					•	-	+		
N75055   2.98   2.09   0.76   1.43   3.93   -		1						•		-	+
## ## ## ## ## ## ## ## ## ## ## ## ##								-		-	-
82905         T69305         1.21         0.48         0.43         2.50         2.85         -         +         -         +           824917         AA489121         1.74         1.05         0.47         1.65         3.69         -         -         -         +         +           950430         AA599078         7.48         3.82         2.21         1.96         3.38         -         -         +         +         +         809657         AA456331         4.72         2.68         1.32         1.76         3.57         -         -         +         +         +         -         -         +         +         -         -         +         +         -         -         +         +         -         -         +         +         +         -         -         -         +         +         +         -         -         +         +         +         +         -         -         +         +         -         -         +         +         -         -         +         +         -         -         +         +         -         -         +         +         -         -         + <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td>-</td><td></td><td></td><td></td></td<>								-			
824917   AA489121   1.74   1.05   0.47   1.65   3.69   + + +   950430   AA599078   7.48   3.82   2.21   1.96   3.38   + + +   950430   AA599078   7.48   3.82   2.21   1.96   3.38   + +   809657   AA456331   4.72   2.68   1.32   1.76   3.57   + +   32509   R43053   4.38   3.20   1.10   1.37   3.97   + + +   346889   W79834   0.89   0.21   0.79   4.21   1.12   + +   145388   R78386   2.66   0.94   1.06   2.83   2.50   - +   4411296   AA682631   1.85   1.07   0.51   1.73   3.59   + + +   726483   AA399269   3.50   2.68   0.87   1.30   4.01   + +   823895   AA490490   1.57   1.72   0.36   0.91   4.40   + +   823895   AA4390490   1.57   1.72   0.36   0.91   4.40   +   845345   AA773478   5.09   3.00   1.41   1.70   3.61   +   133130   R28397   11.12   4.91   3.67   2.27   3.03   + +   135306   AA934762   7.92   5.52   2.06   1.43   3.84   +   1553306   AA934762   7.92   5.52   2.06   1.43   3.84   +   1553306   AA94464531   27.71   20.86   7.03   1.33   3.94   +   1563202   AA01403   9.18   5.49   2.55   1.67   3.59   +   1563202   AA01403   9.18   5.49   2.55   1.67   3.59   +   1563102   AA479133   3.30   1.49   1.08   2.22   3.04   +   15754192   AA479133   3.30   1.49   1.08   2.22   3.04		1						•			
121436 T97457 6.37 6.46 1.46 0.99 4.35 + + 950430 AA599078 7.48 3.82 2.21 1.96 3.38 + + 950430 AA599078 7.48 3.82 2.21 1.96 3.38 + + + 32509 R43053 4.38 3.20 1.10 1.37 3.97 + + + 326889 W79834 0.89 0.21 0.79 4.21 1.12 + + 145388 R78386 2.66 0.94 1.06 2.83 2.50 - + + + 431296 AA682631 1.85 1.07 0.51 1.73 3.59 + + + 46411 H09222 3.07 2.00 0.81 1.54 3.78 + + + 46411 H09222 3.07 2.00 0.81 1.54 3.78 + + + 46411 H09222 3.07 2.00 0.81 1.54 3.78 + + + 46411 H09222 3.07 2.00 0.81 1.54 3.78 + + + 726483 AA399269 3.50 2.68 0.87 1.30 4.01 + + + 726483 AA399269 3.50 2.68 0.87 1.30 4.01 + + + 753092 AA436595 7.06 3.62 2.10 1.95 3.36 + + 1733130 R28397 11.12 4.91 3.67 2.27 3.03 + + + 133130 R28397 11.12 4.91 3.67 2.27 3.03 + + + 1313130 R28397 11.12 4.91 3.67 2.27 3.03 + + + 17867 W90224 13.52 9.26 3.54 1.46 3.82 + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94		1						-			
950430   AA599078   7.48   3.82   2.21   1.96   3.38   -   -   +   +   +   +   +   +   +   +								-			
809657 AA456331		1					1		-		
32509 R43053 4.38 3.20 1.10 1.37 3.97 + + + 346889 W79834 0.89 0.21 0.79 4.21 1.12 + +									-		
346889 W79834 0.89 0.21 0.79 4.21 1.12 + +									-		
145388       R78386       2.66       0.94       1.06       2.83       2.50       -       -       -       -       -       -       -       -       -       -       -       +       +       -       -       -       +       +       +       -       -       -       +       +       +       +       -       -       +       +       +       +       +       -       -       +											
431296       AA682631       1.85       1.07       0.51       1.73       3.59       -       -       + <td< td=""><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td></td><td>_</td></td<>											_
46411       H09222       3.07       2.00       0.81       1.54       3.78       -       -       +       +       +       726483       AA399269       3.50       2.68       0.87       1.30       4.01       -       -       +       +       +       +       823895       AA490490       1.57       1.72       0.36       0.91       4.40       -       -       -       +       +       +       -       -       -       +       +       288695       N59206       3.41       0.88       2.36       3.86       1.44       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       +       +       -       -       -       -       +       +											+
726483       AA399269       3.50       2.68       0.87       1.30       4.01       -       -       +       +         823895       AA490490       1.57       1.72       0.36       0.91       4.40       -       -       -       +         288695       N59206       3.41       0.88       2.36       3.86       1.44       +       +       -       -       -       +         753092       AA436595       7.06       3.62       2.10       1.95       3.36       -       -       -       +         845345       AA773478       5.09       3.00       1.41       1.70       3.61       -       -       -       +         133130       R28397       11.12       4.91       3.67       2.27       3.03       -       -       +       +         788541       AA452955       12.11       3.01       9.65       4.03       1.26       +       +       -       -       +         417867       W90224       13.52       9.26       3.54       1.46       3.82       -       -       +       +         1553306       AA934762       7.92       5.52       2.06		1							-		
823895 AA490490 1.57 1.72 0.36 0.91 4.40 + 288695 N59206 3.41 0.88 2.36 3.86 1.44 + + + 753092 AA436595 7.06 3.62 2.10 1.95 3.36 + 845345 AA773478 5.09 3.00 1.41 1.70 3.61 + 133130 R28397 11.12 4.91 3.67 2.27 3.03 + + + 788541 AA452955 12.11 3.01 9.65 4.03 1.26 + + + 147867 W90224 13.52 9.26 3.54 1.46 3.82 + 611150 AA173109 7.99 6.62 1.96 1.21 4.07 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94								1	_		
288695 N59206 3.41 0.88 2.36 3.86 1.44 + + + +								ŀ	•	•	
753092 AA436595 7.06 3.62 2.10 1.95 3.36 + + 845345 AA773478 5.09 3.00 1.41 1.70 3.61 + + 133130 R28397 11.12 4.91 3.67 2.27 3.03 + + + + 788541 AA452955 12.11 3.01 9.65 4.03 1.26 + + + + 17867 W90224 13.52 9.26 3.54 1.46 3.82 + + + 1553306 AA934762 7.92 5.52 2.06 1.21 4.07 - + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94 + + 1553306 AA934762 7.92 5.52 2.06 7.03 1.33 3.94								j .	+ .		-
845345       AA773478       5.09       3.00       1.41       1.70       3.61       -       -       +       +         133130       R28397       11.12       4.91       3.67       2.27       3.03       -       -       +       +       +         788541       AA452955       12.11       3.01       9.65       4.03       1.26       +       +       -       -       -       +       +       -       -       -       -       +       +       -								1	•	•	+
133130       R28397       11.12       4.91       3.67       2.27       3.03       -       -       +       +       +       -       -       -       +       +       +       -       -       -       -       +       +       -									•	-	+
788541 AA452955 12.11 3.01 9.65 4.03 1.26 + + 4 17867 W90224 13.52 9.26 3.54 1.46 3.82 + + 1 1.553306 AA173109 7.99 6.62 1.96 1.21 4.07 - + + + 1 1.553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1 1.553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1 1.553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1 1.553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1 1.553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 1 1.553306 AA934762 7.92 5.52 1.038 1.62 3.66 + + 1 1.553306 AA934762 7.92 5.55 1.67 3.59 + + 1 1.553306 AA934762 7.92 AA001403 9.18 5.49 2.55 1.67 3.59 + + 1 1.553306 AA934782 AA48821 4.29 1.83 1.47 2.34 2.93 + - + - + 1 1.553306 AA48821 4.29 1.83 1.47 2.34 2.93 + - + - + 1 1.553306 AA479133 3.30 1.49 1.08 2.22 3.04 + + 1 1.553306 AA934762 7.92 AA479133 3.30 1.49 1.08 2.22 3.04 + + 1 1.553306 AA934762 7.92 AA479133 3.30 1.49 1.08 2.22 3.04 + + 1 1.553306 AA934762 7.92 AA479133 3.30 1.49 1.08 2.22 3.04 + + 1 1.553306 AA934762 7.92 AA479133 3.30 1.49 1.08 2.22 3.04 + +								1			+
417867 W90224 13.52 9.26 3.54 1.46 3.82 + + 611150 AA173109 7.99 6.62 1.96 1.21 4.07 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 358456 W96107 37.98 23.51 10.38 1.62 3.66 + 810510 AA464531 27.71 20.86 7.03 1.33 3.94 + 361922 AA001403 9.18 5.49 2.55 1.67 3.59 + 51921 H22928 4.48 1.72 1.68 2.60 2.67 + + - 754192 AA479133 3.30 1.49 1.08 2.22 3.04 + + - 754192 AA479133 3.30 1.49 1.08 2.22 3.04 + + + + + + + + + + + + + +								+	+	•	-
611150 AA173109 7.99 6.62 1.96 1.21 4.07 + + + 1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 358456 W96107 37.98 23.51 10.38 1.62 3.66 + + 810510 AA464531 27.71 20.86 7.03 1.33 3.94 + + 361922 AA001403 9.18 5.49 2.55 1.67 3.59 + 51921 H22928 4.48 1.72 1.68 2.60 2.67 + + - 767994 AA418821 4.29 1.83 1.47 2.34 2.93 + - + - + - 754192 AA479133 3.30 1.49 1.08 2.22 3.04 + +									-	•	+
1553306 AA934762 7.92 5.52 2.06 1.43 3.84 + + 358456 W96107 37.98 23.51 10.38 1.62 3.66 + + 810510 AA464531 27.71 20.86 7.03 1.33 3.94 + + 361922 AA001403 9.18 5.49 2.55 1.67 3.59 + + 51921 H22928 4.48 1.72 1.68 2.60 2.67 + - + - 767994 AA418821 4.29 1.83 1.47 2.34 2.93 + - + - + - 754192 AA479133 3.30 1.49 1.08 2.22 3.04 + +									-	+	+
358456 W96107 37.98 23.51 10.38 1.62 3.66 + + 810510 AA464531 27.71 20.86 7.03 1.33 3.94 + + + + + + + + + + + + + + + + +									-	-	+
810510     AA464531     27.71     20.86     7.03     1.33     3.94     -     -     -     +       361922     AA001403     9.18     5.49     2.55     1.67     3.59     -     -     -     +       51921     H22928     4.48     1.72     1.68     2.60     2.67     -     -     +     -       767994     AA418821     4.29     1.83     1.47     2.34     2.93     +     -     +     -       754192     AA479133     3.30     1.49     1.08     2.22     3.04     -     -     -     +									-	•	+
361922 AA001403 9.18 5.49 2.55 1.67 3.59 + 51921 H22928 4.48 1.72 1.68 2.60 2.67 + - 767994 AA418821 4.29 1.83 1.47 2.34 2.93 + - + - 754192 AA479133 3.30 1.49 1.08 2.22 3.04 +									-	•	
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767994 AA418821 4.29 1.83 1.47 2.34 2.93 + - + - + - 754192 AA479133 3.30 1.49 1.08 2.22 3.04 +									•	+	-
754192 AA479133 3.30 1.49 1.08 2.22 3.04 +									-	+	-
									•	•	
	2543	310 N8115			3.51	2.33	2.92	2  +	•	+	+

Page 10

635110	AA994811	20.00	_		Table	1-3				
121997	T98253				2.23	3.0	n 1			
51011	H19242	2.10			3.36					
283034	NS 1280	7.07			1.88			+	-	
564537	AA121732				3.15		_ I	-	+.	+
827341	777661	1.79			0.93			+	-	•
1631682	AA994801	7.21		1.72	1.05			-	+	+
290378	N64508	1.61	0.53	0.74	3.03	2.19		-	+	+
262865	HOOKEN	17.10 30.48	17.02	4.07	1.00	4.20		+		•
103 1698	A609556	6.26	11.78	11.64	2.59	2.62	. I	-	4	+
109200	T81107	3.17	3.13	1.95	2.00	3.21		+	-	+
824358 A	A489681	0.96	0.93	1.78	3.42	1.79		• .	-	+
326841	R20424	24.32	0.68	0.25	1.42	3.78		+	•	-
240430	H78007	3.52	17.98	6.32	1.35	3.85		•	+	. +
1475200 A	A85830A	28.16	1.28	1.43	2.74	2.46		•	+	+
/3 1272 A	A421073	5.02	22.56	7.13	1.25	3.95		-	-	• *
243860	N55330	30.36	2.47	1.59	2.03	3.16		-	+	+
854874 A	A630346	2.22	12.47 0.68	11.02	2.43	2.76		-	•	+
8241241A	A490802	1.03	0.63	1.15	3.26	1.92	+	+	+	+
1435029 A	A857212	1.84	0.67	0.29	1.63	3.55		+	, <del>-</del>	-
951080IA	A620446	2.43	1.63	0.75	2.74	2.44	1 .	+	-	+
22731	R45264	4.46	2.73	0.66	1.49	3.69		-	•	•
813637 A	4447742	34.08	17.95	1.26	1.64	3 <b>.54</b>		-	+	+
258521	N40743	3.33	1.39	10.41	1.90	3.27	-	-	+-	+
1602619 A	1988798	135.32	117.37	1.20	2.40	2.77	+	-	+	+
144926	R78559	9.20	3.96	33.70	1.15	4.01		•	+	-
897690 AA	1598758	15.87	13.03	3.24 4.03	2.32	2.84	•	+	-	+
740914 AA	478268	7.20	4.16	2.10	1.22	3.94	•		-	· <del>+</del>
1160558 AA 277660	877347	8.03	3.31	2.10 2.94	1.73	3.42	-	_	+	+
840821	N49405	14.16	5.37	5.65	2.43	2.73	-	•	+	+
840821 AA	486261	17.59	8.78	5.60	2.64	2.51	-	+	•	+
760344 AA	426227	5.89	3.00	1.85	2.00	3.14	-	-	-	+
813187 AA	436721	22.70	10.74	7.51	1.96	3.18	•	-	_	+
685516 AA 503851 AA	202573	3.36	1.79	1.03	2.11	3.02	-	-	_	+
1631762 AI	130042	18.15	5.54	9.79	1.88 3.28	3.26	-	•		+
	025126 767309	46.32	36.08	12.06	1.28	1.85	+	+	-	+
_ 1	R18258	15.62	7.28	5.24	2.15	3.84	•	-	-	+
	769348	7.89	2.84	3.37	2.78	2.98	-	-	-	+
	163668	5.14	2.70	1.60	1.90	2.34	•	+	-	
811842 AAA	42177	3.29	1.15	1.46	2.86	3.21	-	-	-	+
810221 AA4	64700	9.91	4.54	3.39	2.18	2.92	+	+	4	-
243/031 H		3.85	1.37	1.68	2.81	2.30	•	-	-	+
178137 H	47015 -	62.65	37.20	18.34	1.68	3.42	•	+	•	•
815800 AA4	85216	38.41	190.87	61.93	1.25	3.85	•	-	-	+
294259 W	01800	1.22 4.21	1.44	0.29	0.84	4.25	-	-	4	+
248454 N	59626	3.70	2.21	1.32	1.90	3.19	•	-	+	+
504290 AA1	5170K	50.88	1.95	1.16	1.89	3.19	-	•	+	+
731257 AA4		4.45	17.89	22.72	2.84	2.24	+	•	+	•
259350 Na	11226 -	4.43 !2.84	2.31	1.41	1.93	3.15	-	•	+	•
1493390 AA89	4627	3.57	6.64	13.93	3.44	1.64	-	-	4	+
6651271AA10	35601	5.48	1.31	1.53	2.73	2.34	•	+	•	•
3776721AA05	KAAA	2.59	2.72 1.39	1.79	2.02	3.06	<del>-</del>	+	•	+
837953 AA45		0.44	6.85	0.81	1.87	3.20	+	-	<u>-</u>	+
	•		4.03	2.94	1.52	3.55	•	-	+	•
						•		-	+	+

Page 11

				т	able 1-3						
1500324 AA88	25210 /	5.77	1.55	1.89	1.49	3.58	•		•	+	+
				3.93	3.06	2.00	+		+	•	•
594693 AA1				2.44	2.05	3.00	-		-	+	+
868119 AA6			4.58	3.49	2.18	2.87	-		•	-	+
502463 AA1		5.77	2.99	1.85	1.93	3.12	-		-	•	_
t t			1.09	2.69	3.60	1.45	+		+	•	+
		1.72	1.47	0.44	1.17	3.88	•		•	+	
430179 AA0				15.13	1.61	3.43	-		_		+
897153 AA6	576961 1	0.77	5.36	3.55	2.01	3.04	•		_	•	+
25517 R	11888	8.52	4.10	2.87	2.08	2.96	•		•	_	+
	004916	2.15	0.93	0.79	2.31	3.77	_		•	+	+
814018 AA	455714	0.98	0.77	0.26	1.28 3.03	2.01	_		+	_	-
843234 AA		7.02	2.32	3.49	3.03 1.06	3.98	_			-	+
823578 AA		0.56	0.53	0.14	3.50	1.54	_		+	-	•
	R <b>6178</b> 0	1.11	0.32	0.72 0.85	2.88	2.15	_		+	-	-
784830 AA		1.83	0.63	7.92	1.91	3.12	-		-	+	-
		24.70	12.93	2.62	3.43	1.59	4		+	-	-
95 <b>0709</b> AA		4.17	1.22	4.08	1.78	3.24			-	+ '	+
1486260 AA		13.23	7.43	1.10	1.20	3.82			-	-	+
878152 A		4.20	3.49 1.21	1.90	3.07	1.95	١.	-	+	-	-
130773	R22156	3.71	0.70	1.34	3.29	1.73		-	+	-	•
586725 A		2.32	23.00	9.58	1.48	3.54	1	-	-	-	+
511521 A		33.96 3.16	1.64	1.02	1.92	3.10	1	-	-	+	+
295206	N75947	· 5.02	1.92	2.10	2.62	2.39	1	+	-	-	•
281072	N50912	2.04	0.70	0.98	2.91	2.09	1	-	+	-	-
824704 A		10.42	4.91	3.62	2.12	2.88		-	•	-	+
143306	R74169	3.15	1.20	1.33	2.63	2.37	1	-	+	-	-
247609	N58148 H95976	4.24	1.94	1.51	2.18	2.82	:	-	-	-	+
250678	A458943	0.47	0.34	0.13	1.38	3.62		-	•	+	+
8250581	A504284	3.42	1.55	1.23	2.21	2.79		+	•	+	+
197888	R96220	4.97	4.63	1.27	1.07	3.92		-	•	+	
261472	H99035	9.31	3.39	4.15	2.75	2.2		+	-		+
137254	R36523	1.35	0.58	0.51	2.35	2.6		-	+	+	+
52730	H29499	6.16	4.38	1.72	1.41	3.5		•	-	+	+
76252	T59873	11.64	8.74	3.19	1.33	3.6		•	+	•	-
898123	AA598487	4.77	1.81	2.03	2.63	2.3 3.0		-		+	+
	AA485214	14.69	7.77	4.77	1.89	۶.¢ 3.9		-	_	+	+
	AA262559	3.59	3.70	0.90	0.97 3.81	3.3 1.1		+	+	-	-
233464	H77297	14.24	3.73	12.38 3.72	1.01		95	-		+	+
80484		14.69	14.52				25	•	+	•	•
322051		11.58	4.27				04	-	•	-	+
299154		1.37	0.71 5 <i>.</i> 59				06	-	•	+	+
136605		10.58	1.16				.81	-	•	-	+
	AA281504		1.34				.59	-	•	+	+
	AA664105		4.1				.45	-	•	•	+
272576							.43	•	-	•	+
83363	1 AA933888					0 2	.14	+	+	•	+
	0 AA496628					4 3	.50	-	-	+	
	5 AA459679				1 2.7		2.24	-	+	-	-
14822					1 1.9		2.96	•	-	+	-
18815					3 2.3	-	2.64	-	•	+	-
	9 AA47848					15	2.78	+	-	τ	
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561000					Table 1-3	}				
261971	AA088886	3.81	3.31	1.01	1.15	3.78	1 -			
266747	AA608546	3.53	2.28	1.04	1.55	3.39	l _	-	+	+
129146		3.05	1.60	1.01	1.91	3.02			<b>T</b>	-
41940		7.97	4.87	2.42	1.64	3.30		-	• .	+
		10.51	5.11	3.65	2.05	2.88		_	•	+.
530014	AA452113	1.46	0.42	1.00	3.46	1.46	+		•	+
287 <b>528</b>	AA070226	46.55	26.81	14.60	1.74	3.19	l .	_	•	•
		1.98	1.06	0.65	1.87	3.05		•	+	+
1425105	AA287323	8.70	5.76	2.55	1.51	3.41		-	· ·	+
248261	AA857632	2.56	1.69	0.75	1.51	3.40		-	+	+
	N78083 AA620458	1.51	0.70	0.55	2.15	2.76		+	+	+
123755		2.03	1.20	0.63	1.69	3.22	_	-	-	+
	R01304	5.06	2.03	2.10	2.49	2.41	+	+	· +	+
927162	AA147499	2.26	1.01	0.84	2.23	2.68	-	-	· •	+
812077	AA521247 AA464617	3.50	2.11	1.08	1.66	3.24	-	•	+	+
1521077	AA906997	7.62	5.19	2.22	1.47	3.43	-		· +	+
416010		2.08	0.72	1.04	2.90	2.00	-	+	·	
	W85832 AA610071	53.71	32.20	16.66	1.67	3.22	-	-	+	-
272576	AA775857	25.15	13.97	8.15	1.80	3.09	<b>-</b> .	•	+	+
420152	AA010233	11.10	9.66	2.97	1.15	3.74	-	-	•	+
417801		4.46	1.54	2.25	2.90	1.98	-	+		
108265	W88848	17.18	13.47	4.77	1.28	3.60	-	-	+	+
	T70810 AA625574	7.24	5.07	2.10	1.43	3.45	-	-	· -	+
365000	AA025112	29.18	19.75	8.59	1.48	3.40	•	-	+	+
112785		4.93	1.78	2.34	2.76	2.11	-	+	•	_
	T859 <b>58</b> AA447 <b>761</b>	7.75	4.81	2.38	1.61	3.25	• •	•	•	+
810284	AA463932	4.96	1.93	2.17	2.58	2.29	-	+		т -
726488	AA463932 AA399264	2.27	1.27	0.74	1.79	3.07	-	-	+	- +
512116	AA133721	10.10	6.03	3.17	1.68	3.18	-	-	· <u>-</u>	+
815760	AA485018	4.59	1.55	2.41	2.96	1.90	+	-	+	-
50344		2.31	1.17	0.81	1.98	2.87	-	-	•	+
293830	H17002 N98283	0.85	0.39	0.32	2.16	2.69	+	-	+	
213890	H72937	4.43	1.72	1.96	2.58	2.26	+	-	+	_
	AA446885	14.14	7.99	4.60	1.77	3.07	-	•	+	+
36607	R46837	10.62	7.56	3.09	1.41	3.44	-	•	•	+
	AA278195	7.52	6.30	2.06	1.19	3.65	•	-	+	+
40017	R53311	4.12 8.29	2.45	1.31	1.68	3.15	-	•	•	+
	AA253444	4.99	4.13	2.94	2.01	2.82	-	-	-	+
490360	AA122048	5.94	2.55	1.74	1.96	2.87	-	-	+	_
544639	AA074677		3.29	1.97	1.81	3.02	-	-	-	+
489109	AA056484	2.82 7.44	1.94	0.84	1.46	3.37	-	•	-	+
279394	N48701	3.18	5.33	2.17	1.40	3.43	٠.	-	+	+
74223	T48411	5.42	1.23 3.10	1.42	2.59	2.23	+	•	+	_
	AA054643	3.54	3.10 1.49	1.76	1.75	3.07	-	•	-	+
712610	AA281793	1.06	0.58	1.46	2.38	2.43	+	-	+	-
448117	AA702480	3.91		0.35	1.83	2.98	-	-	•	+
50114	H16851	0.83	1.64 0.28	1.61	2.38	2.43	-	+	-	+
1472638	AA872323	7.11	∂.28 3.65	0.44	2.92	1.89	+	+	-	+
841399	AA487552	9.08		2.16	1.63	3.29	-	•	-	+
855799	AA664067	9.00	4.37 6.46	3.33	2.08	2.73	-	•	-	+
810753	AA457737	4.95	6.46 1.68	2.64	1.39	3.41	-	-	+	+
51749	H23049	3.65	1.08	2.67	2.95	1.85	-	+	-	-
361659	W96280	13.43	4.23	1.33	2.05	2.75	-	•	+	•
			7.23	8.34	3.18	1.61	-	+	-	-

				. 7	able 1-3					
252453	H87143	20.12	4.53	5.91	1.38	3.40	•	•	` <b>+</b>	+
133717	R27767	9.46		5.91	3.18	1.60	+	+	•	•
785793 A				6.01	2.65	2.14	•	+	-	•
76196	T59668	2.53		1.29	2.83	1.96	+	+	•	•
322561	W15277		70.00	33.31	1.54	3.24	-	-	-	+
825833 A		1.95	1.08	0.66	1.81	2.97	-	-	•	+
755765 A		4.01	2.20	1.36	1.83	2.96	-	•	-	+
1416086 A		8.24	4.30	2.88	1.92	2.86	-	••	+	+
358083	W95279	2.45	0.80	1.44	3.07	1.71	•	+	-	+
	A447731	84.51	49.31	2 <b>7.59</b>	1.71	3.06	-	-	-	+
	A133166	· 5.35	3.78	1.60	1.42	3.35	-	-	+	+
809876 A	A455126	27.74	19.39	8.32	1.43	3.34	-		-	τ
744933	A625812	2.15	1.01	0.82	2.14	2.63	+	•	+	+
321386	W44852	2.45	1.56	0.77	1.57	3.19	-	-	•	+
253577	H89589	1.48	1.24	0.41	1.19	3.57	-	-	-	+
41174	R56106	4.02	2.38	1.31	1.69	3.07	-	-	+	+
149013	R82299.		5.75	2.30	1.36	3.40	-	<b>-</b>	τ	+
296529	W01011	9.83	7.09	2.92	1.39	3.37	-	-	•	+
	AA017200	5.63	3.33	1.84	1.69	3.06	•	•	-	+
154749	R55619	1.76	1.07	0.57	1.65	3.10	•	-	-	+
197176	R92806	20.79	10.24	7.65	2.03	2.72	•	-	+	+
627428	AA190313	24.22	21.29	6.72	1.14	3.61	•	-	<b>T</b>	,
	AA598561	14.33	5.93	6.17	2.42	2.32	•	+	-	+
590298	AA147837	10.43	6.02	3.47	1.73	3.00	-	-	-	+
	AA042928		5.47	2.99	1.67	3.06	-	-	+	ì
1475797	AA872122	85.21	65.01	24.92	1.31	3.42	-	•	+	-
283196	N54728		2.01	1.34	1.89	2.84	-	•	• +	+
	AA126708		13.11	5.65	1.42	3.30	-	-	+	+
	AA873060		4.83	1.85	1.31	3.41	· -	-	+	+
	AA479154		4.46	2.13	1.53	3.20	-	-		+
325606	W52341		3.27	2.93	2.23	2.49 2.60	1 :	_		+
	AA455917		6.63	5.40	2.11	1.92		+	•	_
3207			0.37	0.54	2.79	3.08	Į.	-	-	+
	0 AA77778		4.62	2.45	1.63 1.50	3.21	1	-	+	+
	0 AA96223		4.29	2.01	2.09	2.62	1	-	•	+
	7 AA66415		4.10	3.27 2.40	1.49	3.20		-	+	+
	2 AA67718	5 7.68	5.14	1.40	3.31	1.39		+	-	•
2661			0.59 24.53	6.55	0.99	3.70		-	•	+
	9 AA98870	24.27	6.70	6.06	2.23	2.46		•	+	+
34272			5.11	1.14	0.85	3.84		-	+	+
59014	45 AA 15620 50 AA 48841		1.90	2.57	2.69	1.99		+	-	-
			1.00	0.22	0.84	3.8		-	-	+
2822			11.94					-	+	+
2031			1.86					-	•	+
403 2041								-	+	•
2041	83 AA4314							-	+	•
1419								-	•	+
3568								+	•	•
3300 4717	266 AA033							. <del>-</del>	•	+
246						9 3.	58 -	. <del>-</del>	•	+
	121 AA4850					0 3.	87 -		•	+
701	532 AA284					1 3.	05		•	+
	659 R61						62	- +	-	-
72										Þ

Page 14

		_								
4500	1				Table 1-3					
460841	AA708189	1.02	0.58	0.35	1.75	2.91				
306771		3.61	1.62	1.49	2.23	2.42	_			:
160485		5.17	2.96	1.78	1.74	2.90	_	•	-	+
950461	AA599107	38.86	34.73	11.02	1.12	3.53	•		<b>+</b> .	+
75886		14.08	9.92	4.36	1.42	3.23		-	+	+
283919		7.56	4.94	2.43	1.53	3.11	•	•	+	+
813271	AA455938	0.96	1.12	0.25	0.85	3.79		-	+	+
823627	AA490561	3.25	1.81	1.14	1.79	2.85	•	•	-	+
242706		5.15	2.96	1.78	1.74	2.90	•	•	•	+
970271	AA775957	3.16	1.23	1.52	2.56	2.07	•	+	•	+
783645	AA446690	6.96	2.63	3.51	2.65	1.98		+	-	-
486459	AA042858	6.24	5.89	1.75	1.06	3.57	•	•	-	-
884346	AA629524	5.38	3.61	1.71	1.49	3.14		•	+	+
26406		6.56	5.37	1.92	1.22	3.41		•	. •	+
1467108	AA883160	4.29	2.67	1.42	1.61	3.02	-	•	•	+
1632161		39.75	38.39	11.08	1.04	3.59	•	•	-	+
950473	AA599120	9.36	3.14	5.71	2.98	1.64	-	+	•	+
796885	AA463200	9.64	6.52	3.08	1.48	3.13	•	т	-	•
854122	AA669183	4.20	2.32	1.50	1.81	2.80	•	•	-	+
811911	AA456271	11.89	6.59	4.25	1.80	2.80	-	•	+	+
383,188	AA074224	2.79	1.08	1.39	2.58	2.02	•	+	•	+
	AA699770	3.42	4.00	0.91	0.85	3.74		-	•	-
324342	W47576	4.41	4.29	1.24	1.03	3.57			+	+
470114	AA029295	3.09	1.27	1.44	2.44	2.15	•	+	-	+
	AA489343	28.44	25.71	8.17	1.11	3.48	•	7	-	+
129725	R16957	17.94	10.53	6.22	1.70	2.88		<b>-</b>	-	+
121196		22.80	22.77	6.36	1.00	3.58	-	_	-	+
838668	AA457235	3.58	4.32	0.95	0.83	3.76	-	-	+	+
283142		3.58	1.94	1.31	1.84	2.74	•	_	7	+
1636707	AI017703	19.22	12.06	6.43	1.59	2.99	_	_	•	+
123980	R01637	2.38	0.92	1.19	2.59	1.99	-	+	-	+
253009		115.86	35.75	86.54	3.24	1.34	+	· +		-
461489	AA705058	0.92	0.61	0.30	1.51	3.07	-	_	-	<u>-</u> +
856115	AA630584	1.30	0.82	0.43	1.57	3.00	-		+	+
51151		3.26	3.41	0.90	0.96	3.61	-		+	+
119403		1.22	1.22	0.34	1.00	3.57	-	_		+
	AA455931	4.14	2.24	1.53	1.85	2.71	-	-	_	+
211206		4.71	2.93	1.60	1.61	2.95	_		_	+
127251	AA412053	34.32	29.87	10.07	1.15	3.41	-	_	-	+
627600	AA873577	35.19	14.23	16.93	2.47	2.08	-	+	-	+
02/088	AA197344	7.37	3.62	2.93	2.03	2.52	+	-	+	_
289288	N68998	5.44	3.58	1.79	1.52	3.03	•		•	+
810725	AA457717		1.79	1.20	1.83	2.72	-	_	+	_
243068		9.34	5.76	3.20	1.62	2.92	-	•	+	_
709920	AA430615	9.93	5.82	3.51	1.71	2.83	-	-	+	_
289818		7.47	5.73	2.31	1.30	3.23	•	•	•	+
196345		122.93	41.22	79.12	2.98	1.55	-	+	-	-
74511	AA465424	8.19	4.01	3.29	2.04	2.49	-	-	+	
74511		5.23	3 <b>.37</b>	1.75	1.55	2.98	-	-		+
730313	AA412643	3.95	2.39	1.37	1.65	2.88	-	-	-	• • • · ·
34869		2.74	2.06	0.85	1.33	3.20	•	-	+	
77805	AA608514	28.90	11.69	14.05	2.47	2.0€	-	+	-	•
		15.55	9.75	5.31	1.59	2.93	•	-	-	+
1472304	AA894577	4.23	2.66	1.44	1.59	2.93	•	-	•	+
						•				

Page 15

				_			1				
				. 7	Table 1-3					_	
263070	N20052	1.17		0.37	1.31	3.21	•	•	•	+	
	AA620565	2.02		0.67	1.50	3.02	•	- +	-	<b>T</b>	
308726	N93236	14.73	-	7.73	2.61	1.91	+	<del>1</del>	•	•	
40136	R53348	5.23	1.99	2.77	2.62	1.89	-	-	+	+	
	AA883711	3.95	2.84	1.27	1.39 2.66	3.12 1.85	-	+	•	•	
	AA424344	2.93	1.10	1.58 1.36	1.99	2.52	_	·	•	+	
	AA010223	3.43	1.72	8.14	1.47	3.03		-	-	+	
	AA669443	24.71	16.78 0.90	0.67	1.92	2.59	-	•	• ,	+	
	AA625890 H54020	1.73 · 12.77	10.17	3.93	1.26	3.25	-	-	-	+	
202904 257382	N30699	12.77	10.78	3.86	1.19	3.32	-	-	+	+	
	AA157150	6.42	3.41	2.45	1.88	2.62		-	-	+	
	AA708152	1.30	0.65	0.52	2.00	2.50	•	• '`	-	+	
	AA481135	0.88	0.71	0.27	1.25	3.25	-	-	•	+	
214884	ľ	3.55	2.09	1.27	1.70	2.79	•	-	-	+	
	AA431836	35 <b>.89</b>	20.99	12.92	1.71	2.78	-	-	-	+	
277596		2.91	1.45	1.17	2.00	2.49	-	-	+	-	
344432	1	15.48	11.50	4.93	1.35	3.14	•	-	•	· +	
	AA459750	7.16	4.60	2.44	1.56	2.93	•	-	•	+	
	AA416759	4.84	1.70	2.96	2.85	1.64	•	+	•	-	
	8 AA452566	13.97	9.65	4.61	1.45	3.03	-	-	-	+	
20162		7.51	5.63	2.39	1.33	3.14	-	-	+	+	
	2 AA872057	35.80	21.64	12.69	1.65	2.82	-	•	-	+	
	6 AA598595	1.79	2.04	0.50	88.0	3.59	-	-	-	+	
	7 AA497010	15.69	7.46	6.62	2.10	2.37	-	<del>+</del>	-	+	
48982	3 AA099855	33.49	26.22	10.49	1.28	3.19	-	-	+	+	
81382	5 AA453713	0.52	0.34	0.18	1.52	2.95	-	-		*	
78140	1 AA430205	2.67	1.22	1.18	2.19	2.27	-	-	+	-	
149296	67 AA886333	3.68	1.85	1.49	1.99	2.47	+	+	-	_	
3122	•	4.76	1.64	3.09	2.90	1.54	\	-	_	+	
358		2.91	1.90	1.00	1.53	2.91	1	-		+	
	50 AA599094	1.77	1.02	0.66	1.75 1.70	2.70 2.74	1	-	_	+	
	45 AA465116	4.97	2.92	1.82	1.70	2.48		_	_	+	
	05 AA453926	13.21	6.75	5.32 3.09	1.36	3.08	1 .	-	+	+	
	91 AA452933	9.50	6.97	4.99	1.44	3.00	l	-		+	
	40 AA486516	14.94	10.35 30.51	19.51	1.73	2.71		•	-	+	
	11 AA456975	52. <b>82</b> 7. <b>20</b>	6.82	2.13	1.06	3.38		-	+	+	
	103 AA670353 338 AA775223		2.14	1.14	1.54	2.90	3	-	+	•	
222:			2.10	0.94	1.37	3.05		-	-	+	
	767 AA486300		1.48	0.73	1.46	2.96	;	•	-	+	
907	670 AA496800		2.19	4.18	2.90	1.52	2 -	+	-	-	_
	581 AA400247		1.08	0.36	1.11	3.31	1 -	-	-	+	
	338 H2327		1.24	1.41	2.35			+	-	+	
	398 AA14803			4.63	1.77	2.6	5 -	•	•	+	
	620 AA48803			4.92	1.61			-	. •	+	
	899 AA63037			6.24				-	•	+	
	3253 R3594		1.95	0.94				-	-	+	
	2369 AA15694							<del>-</del>	-		
	3609 W6946		0.58					+	-	+	)
	1496 AA28691	4 2.28					1	-	+	+	
	3252 AA88687	0 9.89						-	+	•	
20	3350 H5436	54 1.9						+	+	+	
45	1707 AA7076	59 7.59	9 5.80	2.43	5 1.3	ı 3.	09   -	•	•	·	

Page 16

					Table 1-	3				
22747		37.20	31.66	11.55	1.18					
815072	AA465180	3.49	1.97	1.33	1.77	3.22	-	_		
251731		0.82	0.84	0.24	0.98	2.62 3.41	1	•	. +	•
815563	AA456833	4.86	3.88	1.55	1.25	3.13	•	<b>.</b>	•.	+
027273	AA190747	1.58	1.06	0.55	1.49	2.88	•	•	•	+
39821	R53428	5.84	5.87	1.73	0.99	3.38		-	-	+
46262	AA598597	8.13	8.02	2.42	1.01	3.35		-	-	+
46367	H09959	1.63	0.62	0.94	2.62	1.74	+	` <b>-</b>	•	+
28436	R40663	0.89	0.37	0.46	2.43	1.93	+	•	•	•
2599 <b>53</b> 307882	N32594	1.46	1.97	0.40	0.74	3.63		•	-	•
717796	W21373 AA283029	7.09	4.18	2.66	1.69	2.67		-	-	+
195903	AA283029	0.89	0.56	0.32	1.58	2.78		_	-	+
42076	R92227	6.95	5.56	2.24	1.25	3.11	١.	_	· . <del>-</del>	+
	R60847 <b>A</b> A887201	5.72	3.44	2.12	1.66	2.69			•	+
201336	AA 128297	2.03	1.50	0.68	1.36	3.00		_	-	+
838470	AA128297 AA457517	51.40	43.89	16.15	1.17	3.18		_	-	+
281191	N50983	2.51	1.48	0.94	1.69	2.66			•	+
391984	A1003625	7.36	3.32	3.45	2.22	2.13	_	+	. <b>-</b>	+
1635665	A1013625	4.56	2.74	1.70	1.66	2.68			-	+
49485		5.13	4.14	1.65	1.24	3.11	-	_	+	+
	H16600 . AA033973	0.54	0.37	0.19	1.46	2.89	_	_	т-	+
427772	AA043347	5.35	3.27	1.98	1.63	2.71	_	_	-	+
701400	AA287948	29.51	20.98	10.06	1.41	2.93	_	_	-	+
144977	R78 <b>725</b>	1.54	1.24	0.50	1.23	3.10	-	_		+
240961	H91011	71.81	48.67	25. <b>09</b>	1.48	2.86	_		-	+
	AA7786 <b>7</b> 5	2.11	0.82	1.20	2.57	1.76	-	+	-	+,
45515	H08417	3.07	1.85	1.15	1.66	2.67	-	<u>.</u>	_	•
	AA446743	3.88	2.73	1.33	1.42	2.91	-	•	+	++
796774	AA443152	0.26	0.10	0.15	2.61	1.72	-	+		- T
415978	W86378	0.44 24.58	0.47	0.13	0.93	3.40	-	-	-	+
68636	T49801	4.83	18.77	8.16	1.31	3.01	-	-	-	+
35812	R14631	2.29	2.97	1.80	1.63	2.69	•	-		+
	AA 129297	1.90	2.07	0.71	1.11	3.21	-	-	-	+
503097	AA151486	7.27	1.12	0.72	1.69	2.63	-	-	_	+
139113	R63022	4.32	4.27 3.44	2.78	1.70	2.61	-	-	+	
131621	R23790	10.21	3.63	1.41	1.26	3.06	•	-	+	+
156386	R73545	2.14	1.88	6.81	2.82	1.50	•	4	-	•
784264	AA447021	5.67	3.30	0.67	1.14	3.18	•	-	-	+
291706	N73480	2.41	1.00	2.18 1 <i>.</i> 27	1.72	2.60	•	•	-	+
310493	W31074	7.56	7.25	2.32	2.41	1.90	+	-	+	•
784016	AA443698	33.80	23.65	11.75	1.04	3.26	•		-	+
282144	N51883	30.16	18.11	11.43	1.43	2.88	-	-	-	+
435341	AA700736	1.73	1.11	0.63	1.67	2.64	-	•	+	•
252904	H88485	5.40	2.26	2.83	1.56	2.74	•	-	•	+
433573	AA701655	10.67	9.66	3.34	2.39	1.91	-	+	•	-
743041	AA406040	6.48	3.98	2.43	1.11 1.63	3.19	-	-	•	+
897971	AA598868	11.55	7.34	4.24	1.63	2.67	-	•	•	+
827168	AA521292	6.51	2.85	3.24	2.29	2.73	•	•	•	+
141361	R64454	1.28	0.68	0.54	1.90	2.01	+	•	+	•
767298	A418383	19.37	16.76	6.18	1.16	2.40	•	•	+	-
436106	A701981	4.14	2.42	1.60	1.71	3.14	•	•	•	+
884388	AA629554	4.89	3.97	1.60	1.71	2.58	•	•	•	+
46294	H09132	1.03	0.35	0.78	2.97	3.06	-	•	•	+
				<del>-</del>		1.311	+	+		_

	_		Ta	ble 1-3					
253253 H89104	1.73	1.30	0.58 1	1.33	2.95	-			
1031346 AA609106	3.21	• • •			2.76	-	-	•	+
429574 AA011445	3.77	1.23	3.11	3.06	1.21	-	+	-	• _
52327 H23459	5.70			1.19	3.08	-	-	-	+
810741 AA457725	72.43			1.50	2.78	•	-	<del>-</del>	Τ
155838 R72215	2.53	1.96		1.29	2.97	-	•	+	+
135083 R33917		17.59		0.97	3.29	•	-	-	+
271472 N35020		7.84	4.28	1.50	2.76	•	-	<u>-</u>	+
29964 R14699		0.63	0.45	1.79	2.47	-	•	_	+
269300 N35863		7.57	3.73	1.41	2.85	•	+	_	_
429678 AA011593		0.86	4.85	3.61	0.64	+	•	-	+
898096 AA598795		2.82	1.45	1.44	2.81	-	• .		+
730554 AA43603		8.38	4.20	1.42	2.83	•	-		+
260628 H9756		7.49	2.93	1.19	3.05	-	_	_	+
730412 AA46996	6 1.25	1.00	0.42	1.25	3.00	•	-	+	_
825771 AA49131	2 12.13	9.01	4.19	1.35	2.90	-	_	-	4
342211 W6378		1.55	0.64	1.24	3.00	-		-	+
845609 AA64434	5 15.29	12.54	5.07	1.22	3.02 2.57	-	-	+	-
814154 AA49625		1.38	0.89	1.67	2.73	-	-	-	+
897730 AA59898	37 41.12	27.32	15.07	1.50	2.18	+		+	
141336 R6382		1.03	0.97	2.05	3.13		•	-	+
263894 N285		0.42	0.15	1.10	3.03	_		•	+
252864 H884		1.00	0.40	1.20 2.18	2.04	•	+	-	•
291057 N721		0.61	0.66	1.55	2.67	-	-	-	+
752690 AA4178		3.90	2.27	2.29	1.93		+	-	+
132911 R261		22.96	27.26	1.34	2.87	_	-	-	+
489663 AA0994	.04 2.11	1.57	0.73 1.30	2.39	1.83	-	+	-	-
430614 AA6780	2.38	1.00	1.30	1.09	3.12	-	-	-	+
1415981 AA9479		3.78	0.14	1.04	3.16		•	~	+
810027 AA4649		0.43	1.45	1.33	2.87	-	<b>-</b> ,	-	+
768469 AA495		3.13 3.13	1.53	1.38	2.82	-	-	-	4.
271076 N42			1.39	1.27	2.91		-	+	•
759948 AA424			2.16	1.52	2.66		-	-	+
529861 AA070			0.87	0.96	3.23	-	-	-	+
838628 AA457 590853 AA161			0.21	1.04	3.14	- 1	-	-	+
	5816 3.19		1.02	1.04	3.14		-	-	+
36387 R46 950464 AA599			3.98	1.72	2.45		-	+	+
898227 AA59			0.68	1.22	2.9		-	-	+
811757 AA46			0.61	1.57	2.5		-	-	+
	5369 6.8		2.95	1.84	2.3		•	-	+
	5285 0.4			1.13			-	_	+
461488 AA70			0.19	1.27			•	_	+
1593502 AA98		25 1.64		1.37			•	+	+
701402 AA21	37936 4.3			1.20					+
	93191 3.			1.55				•	+
251645 H	97033 0.5	50 0.4					·	-	+
50892 H		95 7.4				08			+
826335 AA5	21108 2.	24 1.6				77	- -	+	+
302955 N	91117 10.							•	+
1475738 AA8	72704 238.						_	•	+
884894 AA	69452 41	.65 28.6				.70 .53		-	+
1638479 AIG	16074 3	.27 2.0				.36	•	+	-
20 <b>2154</b>	152503 29	1.46	52 12.40	6 1.7	10 2	۱ ۵۳۰			

Page 18

	_	_			Table 1-3	1				
197821		5.95	4.32	2.15	1.38	2.77	1 _			
951022	AA620421	3.02	1.90	1.18	1.59	2.55				
856535	AA633577	1.40	0.50	1.04	2.79	1.35		•	-	+
841140	AA487020	2.47	3.20	0.73	0.77	3.36	_	+		-
197765	R93767	8.83	3.78	4.92	2,34	1.79	+	-	-	+
125308		2.61	0.93	1.97	2.80.	1.33		•	-	-
323611	W44508	1.25	0.50	0.76	2.48	1.64	•	+	•	-
853998	AA668897	12.39	4.81	8.02	2.58	1.55	-	+	-	•
884766	AA629591	9.95	9.24	3.27	1.08	3.04	-	+	•	•
813712	AA453849	7.62	7.57	2.45	1.01	3.11	-	•	+	+
81 1097	AA485675	2.59	1.28	1.23	2.02	2.10	+	-	+	+
243321	H95088	2.79	1.90	1.05	1.47	2.65	•	•	+	•
856575	AA633658	2.75	2.77	0.88	0.99	3.12	-	•	+	•
360232	AA013099	2.20	1.94	0.74	1.13	2.98	-	• .	+	+
26185	R20616	1.15	0.80	0.43	1.43	2.69	-	•	•	+
469685	AA027912	4.95	4.33	1.67	1.14	2.97	<u> </u>	•	-	+
841501	AA487382	5.94	4.80	2.07	1.24	2.88	-	•	-	+
428215	AA001745	42.11	30.60	15.40	1.38	2.73	•	•	•	+
825356	AA504492	1.18	0.90	0.42	1.32	2.79	•	•	. <del>-</del>	+
884480	AA629719	11.68	10.42	3.91	1.12	2.99	-	•	-	+
813179	AA456717	6.60	2.40	4.88	2.75	1.35	-	•	-	+
645368	AA206267	8.48	7.77	2.82	1.09	3.01	+	-	-	•
	AA232645	0.85	1.07	0.26	0.79	3.30	-	-	-	+
768179	AA426629	3.80	3.11	1.32	1.22	2.87	-	-	-	+
1420830	AA826324	3.06	2.56	1.06	1.19	2.90	-	-	-	+
246541	N77514	10.77	8.71	3.78	1.24	2.85	. <del>-</del>	-	-	. +
742695	AA401521	1.83	1.55	0.63	1.18	2.90	-	•	•	+
451769	AA706795	5.25	5.71	1.66	0.92	3.16	-	-	-	+
827132	AA521232	6.63	4.13	2.68	1.60	2.48	-	•	+	+
460218	AA677457	2.10	1.12	0.95	1.87	2.21	-	•	-	+
40075	R52015	3.13	3.07	1.03	1.02	3.05	_	•	•	+
127928	R08932	3.67	2.74	1.35	1.34	2.73		•	-	+
279785		0.91	0.66	0.34	1.38	2.68	_	•	•	+
1591622	AA983467	49.35	42.75	16.96	1.15	2.91	_	•	+	-
824640	AA482150	6.96	6.37	2.35	1.09	2.97	_	•	-	+
1635350	Al014387	10.17	8.91	3.48	1.14	2.92		•	-	+
855723	AA663960	6.73	7.60	2.12	0.89	3.17		•	+	-
	AA873762	12.48	10.32	4.39	1.21	2.84		•	-	+
845502	A1791165	3.87	3.96	1.26	0.98	3.07	_	•	-	+
897983	AA598874	26.99	22.87	9.42	1.18	2.86	_	_	•	+
824066	AA490734	4.08	3.05	1.51	1.34	2.70	_		-	+
627687	AA196287	56.19	60.68	18.11	0.93	3.10	_		+	•
32887		3.04	2.20	1.15	1.39	2.64	•	_	-	+
433350	AA700604	28.82	38.07	8.83	0.76	3.26		_	+	+
812989	AA464623	2.18	1.75	0.79	1.25	2.77	_		•	-
244154		4.94	3.91	1.79	1.26	2.76	_	-	-	+
278687		13.10	9.87	4.88	1.33	2.69	_	-	-	+
322537		3.18	2.15	1.26	1.48	2.53		-	+	+
399532	AA733038	6.74	4.35	2.74	1.55	2.46	_	-	<del>-</del>	-
854570	AA669124	4.11	2.43	1.78	1.69	2.31		_	•	+
1411726	AA856556	30.68	28.07	10.55	1.09	2.91	_	-	•	+
	AA872153	14.80	6.17	9.24	2.40	1.60	_	+	-	+
80764		26.16	26.33	8.71	0.99	3.00	_			<u>.</u>
825012	AA489194	4.46	3.05	1.76	1.46	2.53		-	•	+
					•		_	-	-	~

S1981   H23422   19.62   13.04   7.89   1.50   2.49						Table 1-3					
159126    AAJSSOOT   3.18   2.86   1.11   1.11   2.88   -	51981	H23422	19.62	13.04	7.89	1.50	2.49	-			
SSEACES  AASSSSSES   20.16								-	-	•	+
S11459   AA115310   37.04   46.42   11.64   0.80   3.18   -   -   +					7.03	1.11	2.87	-	-	•	+
296429   W00890   5.99   4.85   2.18   1.23   2.75   -					11.64	0.80	3.18	-	-	<del>-</del> .	. +
SI3149			5.99	4.85	2.18	1.23	2.75	-	•	•	+
741988 AAG2915	754449 AJ	4410480	1.78	1.58				•	-	•	+
## ## ## ## ## ## ## ## ## ## ## ## ##	813149 A	A456298	6.11					-	-	+	
1492412   AA878561   236.26   205.84   83.73   1.15   2.82   -	741988 A	A402915						-	-	•	
1422794	825842 A	A 504772						-	-	+	
131104								•	-	-	
80292 T44433 2.18 1.64 0.83 1.33 2.62 + + 998844 AA663826 9.12 11.76 2.88 0.78 3.17 + + + + + + + + + + + + + + + + +								-	-	•	
909324   AA663326   912   11.76   2.88   0.78   3.17										•	
750597 21.82 10.94 11.17 2.00 1.95 - + - + - + 750597 21.82 10.94 11.17 2.00 1.75 - + - + - + - + - + - + - + - + - + -								-		•	
303199 W19519 2.17 1.22 1.00 1.77 2.17 - + - + - + + + + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + - + + + - + + + - + + + - + + + + - + + + + + + - +								-	•	+	
## ## ## ## ## ## ## ## ## ## ## ## ##								•	+	-	+
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1471829   AA873351   99.54   79.29   37.33   1.26   2.67   -   -   -   -   -   -   -   -   -							1	•	-	-	
461955 AA779968 1.94 0.91 1.09 2.13 1.78	1							•	•		7
27072								•	_	<b>T</b>	-
282895         N51226         3.91         1.96         2.03         1.99         1.93         +         -         +         -         866866         AA679341         3.00         2.04         1.23         1.47         2.45         -         -         +         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         +         -         +         -         +         -         +         -         +         -         +         -         +         -         +         -         -         +         +         -         +         -         -         -         +         +         -         -         -         +         +         -         -         -         -         +         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -         -								•	т	•	<u>-</u>
866866       AA679341       3.00       2.04       1.23       1.47       2.45       -       -       +         878373       AA670330       31.06       35.83       10.19       0.87       3.05       -       -       +       +         471641       AA034945       4.79       2.74       2.22       1.75       2.16       -       +       +       +       +       +       +       4       4       43680       AA034945       4.79       2.74       2.22       1.75       2.16       -       -       +       +       +       +       +       +       +       4       42366       1.83       1.65       0.68       1.06       2.84       -       -       -       +       +       246800       NS9078       6.67       7.15       2.25       0.93       2.96       -       -       -       +       +       246800       NS9078       6.67       7.15       2.25       0.93       2.96       -       -       -       +       +       256000       -       -       +       4       236       1.13       1.05       1.24       2.64       -       -       -       +       -       -								•	•	<u>.</u>	_
878373 AA670330 31.06 35.83 10.19 0.87 3.05 + + 471641 AA034945 4.79 2.74 2.22 1.75 2.16 - + - + + 47568 AA702422 9.00 7.57 3.31 1.19 2.72 + + 2486076 AA663160 1.94 1.83 0.68 1.06 2.84 + + 248600 N59078 6.67 7.15 2.25 0.93 2.96 + + 565062 AA133896 1.38 1.65 0.45 0.84 3.05 + + 2313 R60946 1.33 1.07 0.51 1.24 2.64 + + 234036 H68988 4.77 3.83 1.81 1.24 2.64 + + 248036 H68988 4.77 3.83 1.81 1.24 2.64 + + 248033 AA629528 4.10 2.83 1.69 1.45 2.43 + + 248037 AA403072 2.75 2.38 1.01 1.16 2.72 + + 25738 AA460286 2.06 1.70 0.77 1.21 2.66 + + 25738 AA660286 2.06 1.70 0.77 1.21 2.66 + + 25738 AA60286 2.06 1.70 0.77 1.21 2.66 + + 25738 AA60286 2.06 1.70 0.77 1.21 2.66 + + 25738 AA60286 2.06 1.70 0.77 1.21 2.66 + + 25738 AA60286 1.67 1.29 0.65 1.29 2.58 + + 25859 AA504784 1.64 1.22 0.65 1.35 2.52 + + 25859 AA504784 1.64 1.22 0.65 1.35 2.52 + + 25859 AA504784 1.64 1.22 0.65 1.35 2.52 + + 25859 AA504784 1.64 1.22 0.65 1.35 2.52 + + 2586340 AA044123 16.47 11.82 6.73 1.39 2.82 + + 2586340 AA044123 16.47 11.82 6.73 1.39 2.45 + + 2586340 AA0362 1.76 2.10 0.59 0.84 2.99 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19 + + 2586459 AA521083 12.36 7.59 5.65 1.63 2.19									-	_	- -
878313       AA00034945       4.79       2.74       2.22       1.75       2.16       -       +       +       +       447568       AA702422       9.00       7.57       3.31       1.19       2.72       -       -       +       +       426800       N59078       6.67       7.15       2.25       0.93       2.96       -       -       +       +       +       565062       AA133896       1.38       1.65       0.45       0.84       3.05       -       -       +       +       +       565062       AA133896       1.38       1.65       0.45       0.84       3.05       -       -       +       +       42313       R60946       1.33       1.07       0.51       1.24       2.64       -       -       -       +       +       42313       AA479100       6.49       5.74       2.36       1.13       2.75       -       -       +       +       424306       H88988       4.77       3.83       1.81       1.24       2.64       -       -       -       +       +       4884333       AA629528       4.10       2.83       1.69       1.45       2.43       -       -       -       +       4       4863403 </td <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td></td> <td>_</td> <td>-</td> <td></td>									_	-	
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234036       H68988       4.77       3.83       1.81       1.24       2.64       -       -       +         884333       AA629528       4.10       2.83       1.69       1.45       2.43       -       -       +         758347       AA403072       2.75       2.38       1.01       1.16       2.72       -       -       +         795738       AA460286       2.06       1.70       0.77       1.21       2.66       -       -       +         72616       TS1995       1.67       1.29       0.65       1.29       2.58       -       -       +         251753       H96908       2.29       3.06       0.74       0.75       3.12       -       -       +         950607       AA608548       19.79       11.09       9.50       1.78       2.08       -       -       +       -         825859       AA504784       1.64       1.22       0.65       1.35       2.52       -       -       +       -         897577       AA496881       0.52       0.50       0.18       1.03       2.82       -       -       +         1599489       AA961361								_		-	+
884333       AA629528       4.10       2.83       1.69       1.45       2.43       -       -       +         758347       AA403072       2.75       2.38       1.01       1.16       2.72       -       -       +         795738       AA460286       2.06       1.70       0.77       1.21       2.66       -       -       +         72616       T51995       1.67       1.29       0.65       1.29       2.58       -       -       +         251753       H96908       2.29       3.06       0.74       0.75       3.12       -       -       +         950607       AA608548       19.79       11.09       9.50       1.78       2.08       -       -       +       -         825859       AA504784       1.64       1.22       0.65       1.35       2.52       -       -       +       -         897577       AA496881       0.52       0.50       0.18       1.03       2.82       -       -       +       -         195652       R89403       18.03       15.86       6.65       1.14       2.71       -       -       +         486340       A								-	•	•	+
758347 AA403072 2.75 2.38 1.01 1.16 2.72 + 795738 AA460286 2.06 1.70 0.77 1.21 2.66 + 72616 T51995 1.67 1.29 0.65 1.29 2.58 + 72616 T51995 1.67 1.09 9.50 1.78 2.08 + 72616 T51995 1.09 9.50 1.78 2.08 + 72616 T51995 1.09 1.09 1.09 9.50 1.78 2.08 + 72616 T51995 1.09 1.09 1.09 1.09 1.09 1.09 1.09 1.09								-	-	-	+
795738 AA460286									_	-	+
72616 T51995 1.67 1.29 0.65 1.29 2.58 + + + + + + + + + + + + + + + + +							2.66	-	•	-	+
251753						1.29	2.58	-	•	-	+
950607 AA608548 19.79 11.09 9.50 1.78 2.08 + + + + + + + + + + + +					0.74	0.75	3.12		-	+	•
825859       AA504784       1.64       1.22       0.65       1.35       2.52       -       -       +         161456       H25546       13.35       10.78       5.08       1.24       2.63       -       -       +         897577       AA496881       0.52       0.50       0.18       1.03       2.82       -       -       -       +         1599489       AA961361       5.51       4.50       2.10       1.22       2.62       -       -       -       +         195652       R89403       18.03       15.86       6.65       1.14       2.71       -       -       -       +         431597       AA676422       5.45       2.95       2.73       1.85       2.00       +       -       +       -         486340       AA044123       16.47       11.82       6.73       1.39       2.45       -       -       +       +         742806       AA400362       1.76       2.10       0.59       0.84       2.99       -       -       +         770935       AA433877       86.21       50.78       40.44       1.70       2.13       -       -       - <td< td=""><td></td><td></td><td></td><td></td><td>9.50</td><td>1.78</td><td></td><td>-</td><td>-</td><td>+</td><td>-</td></td<>					9.50	1.78		-	-	+	-
161456       H25546       13.35       10.78       5.08       1.24       2.63       -       +       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       +       -       -       -       +       -       -       -       +       -       -       -       +       -       -       -       -       +       - <td< td=""><td></td><td></td><td></td><td></td><td>0.65</td><td></td><td></td><td>] -</td><td>•</td><td>•</td><td>+</td></td<>					0.65			] -	•	•	+
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193632 R89403 18.03 13.80 0.03 11.14 2.17 431597 AA676422 5.45 2.95 2.73 1.85 2.00 + - + - + - 486340 AA044123 16.47 11.82 6.73 1.39 2.45 + + 742806 AA400362 1.76 2.10 0.59 0.84 2.99 + + 770935 AA433877 86.21 50.78 40.44 1.70 2.13 + + 53165 R16153 0.89 0.87 0.32 1.02 2.80 + + 7807231 N93428 2.04 1.57 0.81 1.30 2.53 + - 5826459 AA521083 12.36 7.59 5.65 1.63 2.19 + + - + + - + + + + + + + + + + +	1599489	AA961361							-	-	
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742806 AA400362		1							-	, +	-
770935 AA433877 86.21 50.78 40.44 1.70 2.13 + 53165 R16153 0.89 0.87 0.32 1.02 2.80 + 307231 N93428 2.04 1.57 0.81 1.30 2.53 + 526459 AA521083 12.36 7.59 5.65 1.63 2.19 + +								•	-	-	
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826459 AAS21083 12.36 7.59 5.65 1.63 2.19 +		•							-		
826459[AA521083 12.36 7.59 5.65 1.63 2.17]									-	<b>T</b>	
0.0000014.4.014009 45.00 50.00 24.75 1.00 //11 * * * * * * * * * * * * * * * * *									<b>-</b>	_	
868308 AA634008 65.99 60.02 24.35 1.10 2.71 -	86830	8 AA63400	8 65.9	9 60.02	24.35	1.10	2.1	' 1 -	-		·

Page 20

					Table 1-	3				
105620	3 AA621031	1.92	0.89	1.16						
48617	5 AA043133	10.39	9.02	3.91	2.15	1.65				
8148		10.87	5.15	6.40	1.15	2.66	-	•	+	
42904	7 AA007509	3.17	1.48	1.90	2.11	1.70	-	+	•.	•
78369	7 AA446839	3.19	2.89	1.18	2.14	1.67	-	+	•	-
784121	AA446759	1.83	1.82	0.65	1.10	2.70	-	•	-	+
253865	N22007	1.68	1.49	0.63	1.00	2.80	-	-	+	
1291658	AA776810	2.05	1.00	1.17	1,13	2.67	-	-	+	+
595604	AA 167269	9.41	8.19	3.56	2.05	1.75	-	+	-	
784129	AA432085	6.53	5.53	2.50	1.15	2.65	-	-	•	+
251452	H97993	3.12	2.79	1.17	1.18	2.61	-	-	+	•
767346	AA418557	4.50	3.43	1.82	1.12	2.67	-	•	+	
138865	R62742	5.42	4.35	2.14	1.31	2.47	-	•	+	-
770983	AA427401	7.24	6.59	2.70	1.25	2.53	-	-	٠ ـ	+
43705		0.35	0.33	0.13	1.10	2.68	-	-	+	-
76005		2.35	1.80	0.95	1.07	2.71	-	-	•	+
156045	R72518	2.01	1.50	0.83	1.31 1.34	2.46	-	•	-	+
784085	AA446639	3.93	3.17	1.56	1.24	2.43	-	•		+
418053		7.57	7.14	2.80	1.24	2.52	-	•	-	+
322192		7.97	9.95	2.69	0.80	2.70	-	•	-	+
309864		2.49	1.62	1.12	1.54	2.96	-	•	+	+
44975		8.91	10.72	3.05	0.83	2.22	+	-	+	-
825719	AA504834	0.76	0.67	0.29	1.14	2.92	-	-	-	+
291394		6.46	4.22	2.92	1.53	2.62 2.22	-	-	•	+
796313	AA461311	0.78	0.80	0.28	0.98		•	•	+	-
810734	AA480820	2.46	1.75	1.05	1.40	2.77	-	•	•	+
488033	AA045792	3.14	2.24	1.34	1.40	2.34	-	-	-	+
884546	AA629808	156.92	121.14	64.25	1.30	2.44	-	-	-	+
41411 247546	R56885	112.58	99.23	43.25	1.13	2.60	-	-	-	+
		11.16	8.98	4.48	1.24	2.49	-	•	•	+
37665	AA496173	3.75	3.51	1.41	1.07	2.67	-	•	+	-
563461	R61374	1.01	0.72	0.43	1.40	2.33	-	-	-	+
775502	AA113347 AA292995	4.17	1.82	2.90	2.29	1.44	-	-	-	+
1522700	AA292995 AA902249	4.55	3.15	1.99	1.45	2.28	_	+	•	•
268178		1.23	1.56	0.42	0.79	2.94	-	•	•	+
757222	N35772 AA496148	28.02	27.47	10.37	1.02	2.70	-	_	• •	+
1475028	AA857413	5.14	3.76	2.18	1.37	2.36		-	+	+
840530	AA487934	249.73	233.03	94.34	1.07	2.65	-	_	T-	-
950989	AA620404	4.05	3.61	1.56	1.12	2.60	_	-	-	+
756556	AA481438	2.48	1.95	1.02	1.28	2.44	•	•	-	+
868396	AA634164	17.53	6.47	17.49	2.71	1.00	+	•	_	+
857681	AA633768	6.77 54 <b>.39</b>	5.15	2.83	1.32	2.39	-	•	+	+
1555523	AA974971	1.21	52.93	20.36	1.03	2.67	-		_	+
206785	R98047	6.05	0.85	0.53	1.41	2.28	-	•	+	· •
781061	AA446479	0.73	5.71	2.30	1.06	2.63	-	-	•	+
345056	W72798	0.73	0.44	0.35	1.64	2.05	-	-	_	4
	AA496736	2.58	0.35	0.71	2.47	1.22	-	+	•	-
376764	AA046321	2.58	3.15	0.90	0.82	2.87	•	•	-	+
452936	AA778869	5.15	2.44	0.98	1.06	2.63	•	-	-	
76605	T51043	2.48	4.43	2.04	1.16	2.52	-	•	-	+
859832	AA668531	2.46 2.57	2.10	0.99	1.18	2.50	•	•	-	+
809995	AA454862	2.80	2. <b>48</b> 2. <b>07</b>	0.97	1.04	2.64	-	•	-	+
632137	AA167728	0.80	2.07 0.57	1.21	1.35	2.32	-	•	-	+
		0.00	4.57	0.35	1.41	2.27	-	-	-	·

					Table 1-3					
122183	T98628	7.61	10.97 .	2.56	0.69	2.98	•			
	N64762	1.10	0.77	0.49	1.42	2.24	•	•	-	+ .
815772 A.A		1.44	1.12	0.61	1.29	2.38	-	-	-	+
629498 A		3.73	2.69	1.64	1.39	2.28	•	-	-	+
	N33927	2.99	2.12	1.33	1.41	2.25	•	-	•	+
838504 A		41.97	41.10	15.95	1.02	2.63	-	-	•	+
450711 A		1.29	1.51	0.46	0.86	2.79	•	•	-	+
83011	T70522	32.97	35.26	12.18	0.93	2.71	•	-	•	+
866882 A		10.73	11.79	3.94	0.91	2.73	-	•	-	+
277761	N49605	2.43	1.65	1.12	1.47	2.17	•	•	•	+
489722 A		4.03	4.49	1.47	0.90	2.74	•	•	+	+
796527 A.		1.35 0.63	1.17	0.55 0.29	1.15 1.47	2.48	•	•	+	-
33045 137984	R19478 R63129	1.63	0.43 1.42	0.66	1.15	2.13	-	<u>.</u> .	. +	-
1574594 A		3.08	2.54	1.28	1.13	2.40		_		+
138861	R62780	1.72	1.35	0.74	1.27	2.33	•	_	-	+
289645	N62866	0.92	0.67	0.42	1.38	2.21	•	-	+	•
140792	R67081	1.66	0.85	1.02	1.96	1.63	+	-		_
130136	R20813	0.29	0.22	0.12	1.29	2.29	-	-	+	-
503741 A		3.64	4.28	1.33	0.85	2.74	-	-	+	+
	A521243	2.52	2.65	0.96	0.95	2.63	•	•	-	+
841348 A	A487650	11.96	14.60	4.33	0.82	2.76	-	-	•	+
949940 A	A599178	12.52	17.40	4.38	0.72	2.86	•	-	-	+
450464 A	A682819	3.97	2.31	2.13	1.72	1.86	-	-	•	+
743211 A	A400214	3.84	1.71	2.87	2.24	1.34	+	•	-	-
1.	A676920	0.80	0.68	0.33	1.17	2.41	•	-	-	+
	AA630 <b>620</b>	0.25	0.18	0.11	1.36	2.21	•	-	-	+
142326	R70649	1.49	0.78	0.90	1.92	1.65	+	-	-	-
1	AA872143	224.84	89.33	214.43	2.52	1.05	+	•	•	-
	AA420992	2.04	1.38	0.98	1.48	2.09	-	-	+	- +
1	AA677499	14.78	16.78	5.51	0.88	2.68 2.63	-	•	-	+
244202	N52973 AA456931	9.25 53.19	9.97 53.66	3.51 20.71	0. <b>93</b> 0. <b>99</b>	2.63 2.57		<u>.</u>	-	+
	AA774230	6.47	4.45	3.08	1.45	2.10	] -	-	-	+
251417	H96504	0.69	0.68	0.27	1.01	2.52			•	+
	AA598675	2.02	3.02	0.71	0.67	2.86	] _	•	-	+
	AA779221	4.95	6.83	1.77	0.72	2.80	1 -	-	+	-
	AA455482	50.59	51.18	19.96	0.99	2.53	1 -	_	-	+
278531	N66158	57.48	74.35	20.99	0.77	2.74		-	•	+
	AA625788	2.18	2.08	0.89	1.05	2.46		•	-	+
251727		4.88	3.87	2.17	1.26	2.25		•	•	+
	AA669451	8.02	8.08	3.19	0.99	2.51		•	, •	+
	AA046489	54.67	58.82	21.22	0.93	2.58		-	•	+
	AA486403	11.11	10.81	4.49	1.03	2.48		•	•	+
	AA418943	3.64	3.03	1.58	1.20	2.30		-	-	+
	AA489478	31.17	29.51	12.76	1.06	2.44		-	•	+
	AA976544	3.11	3.33	1.21	0.93	2.56		- -	-	+
46620		5.25	2.02	6.01	2.61	0.87		+	•	+
	AA630376	5.69	5.26	2.37	1.08	2.40		-	•	• +
	AA970720	2.41	3.27	0.88	0.74	2.73			-	+
	AA454634	3.13	4.22	1.15	0.74	2.73	•	-	-	-
149895	AA487680 H00752		5.15 4.56	11.09 2.84	2.36 1.33	1.10 2.11		-	+	_
	AA682274		1.35		2.55	0.9		-	-	•
401073	1777002214	J.77		2.04	2.55	0.7	- 1 '			

					Table 1-	3				
743574	AA609456	6.42	4.16	201			_			
823850	AA490456	0.78	4.46	3.21	1.44	2.00	-	•	-	+
435330	AA699926	10.36	0.80	0.32	0.98	2.46	-	-	+	
84078	T70999	0.34	11.94 0.26	4.03	0.87	2.57	-	•	+.	
415415		9.68		0.16	1.29	2.14	-	-	-	+
285226		25.74	10.64	3.83	0.91	2.52	~	•	•	+
41207		2.05	40.08	9.23	0.64	2.79	-	-	+	_
122239		6.50	2.42	08.0	0.85	2.57	-	•	-	+
284865		2.87	5.28 2.71	2.98	1.23	2.18	-	•	-	+
	AA780841	5.61	3.47	1.22	1.06	2.34	-	-	-	+
50562		34.20	30.09	3.15	1.61	1.78	-	-	-	+
71672		4.77	4.16	15.14	1.14	2.26	•	•	+	
	AA135870	31.62	13.68	2.13	1.15	2.23	•	•	-	+
361323	AA017544	4.53	6.49	30.14	2.31	1.05	-	+	•	•
111389	T84308	2.59	2.15	1.70	0.70	2.66	-	-	+	-
<b>7</b> 31337	AA416782	0.75	0.75	1.21 0.32	1.20	2.14	-	-	-	+
455271	AA677574	1.64	1.62	0.32	1.01	2.32	-	•	•	+
488301	AA085748	0.97	1.24	0.38	1.01	2.30	-	•	+	•
511107	AA088371	7.14	7.60	3.02	0.78	2.52	-	•	+	-
626462	AA188999	5.86	3.92	3.24	0.94	2.36	-	-		+
897906	AA598652	10.34	13.66	4.08	1.49	1.81	-	•	•	+
<b>810</b> 408	AA457092	9.85	10.74	4.18	0.76 0.92	2.54	-	-	-	+
22600	T87235	0.36	0.18	0.28	1.99	2.36	. •	-	_	+
343688	W69160	6.94	5.30	3.54	1.31	1.29	+	-	-	-
461761	AA682399	8.65	14.46	3.24	0.60	1.96	-	•	+	-
784285	AA447503	3.99	2.02	3.12	1.98	2.67	-	-	-	+
845780	A1791127	5.95	4.75	2.97	1.25	1.28 2.00	-	+	-	-
266500	N22687	7.14	7.39	3.12	0.97	2.29	-	-	-	+
119851	T94611	6.52	3.28	5.16	1.98	1.26	-	-	-	+
809824	AA455519	2.66	3.75	1.05	0.71	2,54	•	+	-	-
782513	AA432030	5. <b>78</b>	4.46	2.98	1.29	1.94	-	•	<del>-</del>	+
048046	AA206914	15.56	14.08	7.32	1.11	2.13	_	•	+	-
704076	AA279172	5.70	2.95	4.40	1.93	1.30	-	+	-	+
6332361	AA782292	2.75	2.53	1.30	.1.08	2.12	_	-	-	-
739070	AA190997	11.04	13.42	4.64	0.82	2.38	-	_	+	•
757101	AA421769	15.96	26.86	6.13	0.59	2.61	-	-	+	•
740700	AA443966 AA477283	6.08	6.70	2.66	0.91	2.29	-	-	+	-
427812	AA477283 AA001614	4.71	4.39	2.24	1.07	2.10	•	•	+	-
1613940	AA001614 AI000294	9.94	12.03	4.27	0.83	2.33	-	-	+	-
	A1000294 AA001444	3.48	3.17	1.70	1.10	2.05	-	•	_	_
	AA486370	2.86	1.43	2.52	2.01	1.14	+	-	-	
220069	H85437	7.45	7.83	3.42	0.95	2.18	-	-	-	+
	AA988630	4.81 13.38	8.77	1.92	0.55	2.50	-	-	_	+
41287	R56613	0.72	13.90	6.58	0.96	2.03		•	+	•
	AA677706	24.08	0.52	0.45	1.38	1.59	-	•	+	-
151477	H02837	2.44	105 <b>.26</b> 2 <b>.74</b>	8.92	0.23	2.70	-	•	+	+
362080	AA001434	3.81	1.72	1.21	0.89	2.01	-	•	+	•
289832	N63172	6.67	4.11	5.5 <b>6</b>	2.21	0.69	•	+	•	-
767784	AA418670	8.86	14.76	5.4 <b>6</b>	1.62	1.22	+	-	•	•
122428	T99280	2.03	1.78	4.05	0.60	2.19	•	•	+	•
753794	AA406115	1.65	0.94	1.29 1.78	1.14	1.57	-	•	+	•
563444	AA112660	6.23	10.09	3.10	1.75 0.62	0.93	. +	•	•	•
740941	AA478298	12.33	29.84	6.23	0.62 0.41	2.01	•	• .	•	+
•				V.2J	V.4 I	1.98	•	. <del>-</del>	+	•

				Table 1-3	3				
462953 AA682423	3.21	5.03	1.90	0.64	1.69	•	-	+	-
1492202 AA875913		1.98	2.70	1.31	0.96	+	•	-	-

Table 1-4

					Table	e 1-4						
~	GenBank	Aver. of	Aver. of	Aver.	Tumor	Tumor	T/N	· •	,			
Clone		Tumors	Normals	of BPH	Normal			EV Sam	T/N	T/B	T/B	T/B
768180	AA453562	25.18	0.81	0.50	31.24	50.69	+	374 -313	3X,>5/5	10X, >1/5 5	X, >3/5	3X,>5/5
1024472	AA196979	24.42	1.02	0.75	23.91	32.70	+	+ '	+	+	+	+
2034473	AA779728	33.61	1.51	1.35	22.27	24.97	+	+	+	+	+	+
281003	N50880	128.77	6.30	4.90	20.43	26.26	+	+		+		
308682 281039	W25202	8.99	0.87	0.27	10.37	32.91	+	+	+	+	+	+
		14.05	0.94	0.90	14.99	15.57	+	+	+	+	+ ·	+
284701	N64840	33.05	7.05	1.52	4.68	21.68	•			+		
623940	AA490213	22.39	1.32	2.83	16.99	7.91	+			+ .	+	+
42415	R60981	6.86	0.51	0.72	13.52	9.49	+	+	+	+		
46166	H09076	13.13	0.93	1.71	14.12	7.68	+	+		+		
784168	AA432103	53.04	7.37	3.83	7.20	13.84	T	+		+		
773479	AA427899	12.32	0.65	5.99	18.85	2.05		+	٠,	. +		+
222157	H85806	8.80	1.02	0.72	8.67	12.14	+	+	+			
486035	AA043109	24.75	3.47	1.86	7.13	13.32	+	+	+	+		+
450049	AA703396	18.37	1.83	1.80	10.05	10.21	+	+	+	+	+	+
66336	T66832	1.67	0.63	0.10	2.63	16.70	+	+	. +	+	+	+
35058	R45192	15.59	1.89	1.44	8.24	10.70				+		
26736	R39891	9.52	3.36	0.62	2.83	15.37	+	+	+	+		+
	AA159714	28.06	5.61	2.29	5.00	12.23				+		+
32160	R43432	26.80	3.59	2.77	7.47	9.69	+			+		+
	AA465495	14.66	1.70	1.75	8.61	4		+	+	+	+	+
292515	N68465	16.22	2.86	1.56	5.67	8.38	+	+		+		
730739	AA436097	36.49	7.10	3.45	5.14	10.38			+	+	+	+
282089	N48259	11.18	1.51	1.41	7.4 <b>1</b>	10.57	+		+	+	+	+
283312	N54763	63.94	6.63	12.14	9.65	7.92	+			+		
291255	N72215	15.20	1.35	4.65	11.29	5.27	+	+	+			
1048810	AA621342	20.15	3.04	2.59	6.63	3.27	+	+	+			
197525	H52119	9.52	1.85	1.06	5.16	7.78	+	+	+	+		+
67401	T49326	1.47	0.20	0.23	7.28	9.01				+		+
490178	AA 121266	24.55	5.63	2.73	4.36	6.27 8.99	+			+		
1556526	AA935560	0.88	0.15	0.12	5.96	7.33	+			+		+
746152	AA419486	9.35	2.23	1.07	4.18	8.70	+			+		
	AA780190	22.40	3.90	3.28	5.74	6.82				+		
259374	N31952	10.51	2.33	1.31	4.50			+	+		+	+
757435	AA437224	365.78	59.86	58.35	6.11	8.04 6.27				+		
754280	AA479284	23.83	3.20	4.92	7.46	4.85	+		+	+		+
877835	AA625634	23.06	4.34	3.34	5.32	6.90	+	+	+			
/12950 /	AA282272	11.53	3.21	1.34	3.59	8.58		+	+			+
50772	H16803	18.10	6.74	1.91	2.68	9.47	+			+		
1636908	A1000271	6.58	4.38	0.63	1.50	10.44				+		+
432194	AA679414	10.34	1.11	4.04	9.35	2.56				+	+	+
488964	AA047260	7.77	2.13	0.96	3.64	8.08	+	+	+			
1412481	AA845156	8.41	3.50	0.90	2.40	9.31	+			+		
283173	N45236	17.85	3.97	2.47	4.49	7.21	+			+		
811867	AA454963	24.57	2.97	7.43	8.28	3.30	+			+		
26519	R20669	36.05	19.25	3.76	1.87	9.59	+	+	+			
687297	AA235224	12.18	3.12	1.63	3.90		٠			+	+	+ '
666334 /	AA232249	1.64	1.54	0.16	1.07	7.48		•			+	+
1468260 A		4.53	1.13	0.65	4.03	10.14				+	+	+
258300	N30680	18.15	4.53	2.65	4.01	6.96				+		
971279	AA682905	54.34	6.63	22.20	8.20	6.86				+		2
1610448	AA991856	135.55	57.06	16.44	2.38	2.45		+	+			
				· • • • • •	20	8.25				+		+

## ## ## ## ## ## ## ## ## ## ## ## ##				Table 1	I <del>-4</del>						
280217 N47941 20.52 5.86 2.89 3.50 7.09 +	461522 AA705237	1.21 0.65	0.14	1.87	8.74				+		
240694 H78135 4.53 0.90 0.83 5.01 5.47 321706 W35398 7.80 1.86 1.25 4.20 6.23 + 770066 AA430545 11.43 2.60 1.93 4.40 5.94 + 811603 Al732784 19.62 2.74 6.22 7.16 3.15 + 51773 H23211 11.48 2.40 2.09 4.77 5.50 + 66317 T66815 5.50 1.76 0.78 3.13 7.06 + 35626 R45292 9.42 1.09 6.51 8.67 1.45 + + 450307 AA682851 9.39 1.62 2.25 5.80 4.17 + 447715 AA702781 14.83 7.21 1.88 2.06 7.90 + 731290 AA416843 11.60 4.56 1.57 2.54 7.40 + 825857 AA504783 4.27 2.51 0.52 1.70 8.20 + 565235 AA136125 35.83 12.59 5.15 2.85 6.96 + 756600 AA481464 47.06 7.13 14.79 6.60 3.18 + + 293635 N63807 26.47 3.44 12.94 7.69 2.05 + + + 757392 AA426110 0.86 0.37 0.12 2.32 7.41 +	868282 AA633957	8.99 2.01	1.47	4.48	6.12			+		+	+
321706 W35398 7.80 1.86 1.25 4.20 6.23	280217 N47941	20.52 5.86	2.89	3.50	7.09	+			#		
770066 AA430545 11.43 2.60 1.93 4.40 5.94 +	240694 H78135	4.53 0.90	0.83	5.01	5.47			+			+
811603       A1732784       19.62       2.74       6.22       7.16       3.15       +       +       +       +       51773       H23211       11.48       2.40       2.09       4.77       5.50       +											
51773       H23211       11.48       2.40       2.09       4.77       5.50       +       +       +       +       +       66317       T66815       5.50       1.76       0.78       3.13       7.06       +						+			+		
66317 T66815 5.50 1.76 0.78 3.13 7.06							+				
35626 R45292 9.42 1.09 6.51 8.67 1.45 + + + + + + + + + + + + + + + + + + +					1			+			+
450307 AA682851 9.39 1.62 2.25 5.80 4.17 + + + 447715 AA702781 14.83 7.21 1.88 2.06 7.90 + + 731290 AA416843 11.60 4.56 1.57 2.54 7.40 + + 825857 AA504783 4.27 2.51 0.52 1.70 8.20 + + + 565235 AA136125 35.83 12.59 5.15 2.85 6.96 + 756600 AA481464 47.06 7.13 14.79 6.60 3.18 + + + 293635 N63807 26.47 3.44 12.94 7.69 2.05 + + + + + 757392 AA426110 0.86 0.37 0.12 2.32 7.41 + +									+		
447715 AA702781       14.83       7.21       1.88       2.06       7.90       +         731290 AA416843       11.60       4.56       1.57       2.54       7.40       +         825857 AA504783       4.27       2.51       0.52       1.70       8.20       +       +         565235 AA136125       35.83       12.59       5.15       2.85       6.96       +       +         756600 AA481464       47.06       7.13       14.79       6.60       3.18       +       +         293635       N63807       26.47       3.44       12.94       7.69       2.05       +       +         757392       AA426110       0.86       0.37       0.12       2.32       7.41       +       +								+			
731290 AA416843					1	+	+				
825857 AA504783											
565235 AA136125 35.83 12.59 5.15 2.85 6.96 + T56600 AA481464 47.06 7.13 14.79 6.60 3.18 + + T56355 N63807 26.47 3.44 12.94 7.69 2.05 + + T57392 AA426110 0.86 0.37 0.12 2.32 7.41 + T57392 AA426110 0.86 0.37 0.12 2.32 7.41					,				+		
756600 AA481464 47.06 7.13 14.79 6.60 3.18 + + + 293635 N63807 26.47 3.44 12.94 7.69 2.05 + + + + + 757392 AA426110 0.86 0.37 0.12 2.32 7.41 +					•					+	+
293635 N63807 26.47 3.44 12.94 7.69 2.05 + + + + + 757392 AA426110 0.86 0.37 0.12 2.32 7.41 +									+		
757392 AA426110 0.86 0.37 0.12 2.32 7.41 +											
						+	. +	+			
											+
	595695 AA173189		0.44	1.54	8.16					+	+
151055 H02230 27.32 7.51 4.62 3.64 5.92 +					1			,	•		
838366 AA457172 11.15 1.82 3.27 6.12 3.41 + + + 460279 AA677629 0.87 0.46 · 0.11 1.87 7.58 + + +							т	Τ		_	
											_
5,027 1411,3350 Sio, 27,0									7	•	•
1631713 AI025015 5.68 0.93 1.77 6.13 3.21 + + + 564801 AA129551 12.35 3.96 1.99 3.12 6.21 + +							Τ	т	_		
· · · · · · · · · · · · · · · · · · ·							4.		•		
						•					
745347 AA625666 2.15 0.29 1.14 7.39 1.88 + + + 772304 AA404486 113.14 18.10 38.24 6.25 2.96 + +					1						
							•	•			+
81331 T60075 2.84 0.78 0.51 3.63 5.53 + +						4			+		•
840894 AA482340 34.81 11.87 5.68 2.93 6.15 +						•			•	+	
839081 AA488732 16.60 5.37 2.78 3.09 5.98 +					•				+		
1					1						+
· · · · · · · · · · · · · · · · · · ·					1						+
								+			+
84955 T74846 5.26 1.01 1.43 5.21 3.69 +							+				
159608 H16152 14.43 4.14 2.69 3.49 5.36 +									+		
1390584 AA843592 6.54 1.91 1.21 3.42 5.41 +					1				+		
825736 AA504842 0.92 0.60 0.13 1.54 7.26 +										+	
				1.51	7.24					+	+
787938 AA452278 5.09 1.27 1.08 4.01 4.71 +			7 · 1.08	4.01	4.71					+	
809603 AA442991 49.03 6.94 29.52 7.06 1.66 + + +	809603 AA442991	49.03 6.9	4 29.52	7.06	1.66	+	+	+			
020107 1021321 0.13 0.34 0.10 1.30 1.21	826109 AA521327	0.75 0.3	4 0.10	1.38	7.27	l					+
884606 AA630006 33.46 13.42 5.43 2.49 6.16 +	884606 AA630006	33.46 13.4		2.49		1			+		
023022 AA71310 14:01 7:40 1:10 1:10	825822 AA491370	14.07 9.	18 1.97	1.48		1			+		+
531028 AA070495 21.14 3.89 6.62 5.44 3.20 + +	531028 AA070495	21.14 3.		5.44		1	+	+			
814062 AA465338	814062 AA465338					1			+	+	
826103 AA521416 1.31 0.82 0.19 1.61 6.93	826103 AA521416					1				+	
461670 AA779919 8.90 1.72 2.71 5.19 3.28 +	461670 AA779919					1		+			
415700 W85697 3.36 0.61 1.15 5.53 2.93 +	415700 W85691					1	+				
815163 AA481144 1.20 0.57 0.19 2.09 6.36	815163 AA481144					1				+	
810558 AA464568 14.49 3.00 4.01 4.83 3.61 +						1		+			
810220 AA464689 2.17 0.39 0.77 5.60 2.82 +							+				-
359781 AA010932 7.82 2.50 1.49 3.13 5.26	359781 AA01093	7.82 2.	50 1.49	3.13	5.26	i					+

					Table	1-4						
95265	AA451807	4.58	1.26	0.97	3.63	4.74				+		
40306	AA485333	2.49	1.26	0.39	1.97	6.38				•	_	
	AA488986	14.11	4.37	2.84	3.23	4.96			•		•	+
86918	H43317	3.04	1.10	0.56	2.76	5.38				+		Ψ.
91904	H38880	7.67	1.44	2.75	5.32	2.79		+	+	•		
71317	AA476234	9.44	1.78	3.46	5.31	2.73			+			
	AA521394	1.20	0.83	0.18	1.44	6.59				+		
30428	R42061	4.29	0.99	1.16	4.35	3.68	+			+	•	
	AA620995	10.33	4.78	1.77	2.16	5.84				+		
51806	H96647	2.94	1.83	0.46	1.60	6.39				·		+
:05538	N89861	11.44	3.31	2.54	3.46	4.50					•	•
52933	H29315	3.59	1.99	0.58	1.80	6.14				•	•	+
	AA490522	1.38	1.09	0.21	1.27	6.65			•		+	•
	AA666366	85.06	37.74	15.20	2.25	5.60					•	+
	AA018675	16.82	3.11	6.91	5.41	2.43			+			•
153132	H89036	1.40	1.21	0.21	1.16	6.66						+
:33419	H77706	5.55	1.14	1.90	4.87	2.93		+				•
	AA953229	5.30	1.51	1.24	3.52	4.28			+			+
	AA608636	3.00	1.87	0.48	1.60	6.18				+		+
	AA620591	6.84	6.68	1.01	1.03	6.75				+ .		·
	AA427395	10.54	2.90	2.58	3.63	4.08						+
	AA486533	79.86	42.51	13.69	1.88	5.83				+		
268211	N36327	3.43	0.70	1.23	4.93	2.78	+					
	AA487914	15.85	7.18	2.88	2.21	5.50						+
	AA461092	30.31	5.27	15.52	5.75	1.95		+	+			
	AA458516	10.13	5.34	1.75	1.90	5.78						+
150262	AA777488	42.54	7.72	19.71	5.51	2.16		+	+			
	H01129 AA449837	13.66	3.31	3.87	4.12	3.53			+			
	AA045699	7.66	2.78	1.58	2.76	4.86						+
	AA136386	39.41	23.94	6.63	1.65	5.95				,		+
	AA054701	3. <b>93</b> 39.47	2.97	0.63	1.32	6.27	•			+		
	AA461467	6.21	7.39	17.54	5.34	2.25		+	+			
	AA465598	12.37	4.35	1.01	1.43	6.16				+		+
53022	R15880	1.34	6.41 0.70	2.19	1.93	5.65					+	
	AA055306	12.60	3.02	0.24	1.90	5.63						+
	AA481045	1.12	0.67	3.78	4.17	3.34	+					
51700	H22853	3.66	1.21	0.19 0.82	1.67	5.84					+	
80050	T63893	8.15	3.84	1.52	3.02	4.47						+
594994	AA172372	14.61	10.98	2.38	2.12	5.35		•		+		
	AA460346	5.39	1.25	1.71	1.33 4.31	6.14 3.15						+
	AA490611	0.59	0.37	0.10	1.58	5.13	+					
	AA100674	225.66	92.92	45.17	2.43	5.00					+	
	AA496957	9.40	3.54	1.97	2.65	4.76				+		
784130	AA432075	17.23	6.93	3.50	2.49	4.93				+		
1535596	AA918370	401.67	285.67	66.91	1.41	6.00						+
1605407	AA988746	9.00	4.61	1.65	1.95	5.45						+
	AA996028	23.76	9.09	4.98	2.61	4.77				•		
796 <b>239</b>	AA460675	26.07	23.22	4.18	1.12	6.24					+	
493 <b>02</b>		4.88	0.86	2.90	5.68	1.68	+			~	+	
	AA436142	12.77	7.46	2.26	1.71	5.64	•				+	
207 <b>029</b>	R98842	81.88	14.79	45.41	5. <b>54</b>	1.80		+	+		*	
	AA458747	1.34	1.36	0.21	0.99	6.35		•	•			
1636108	AI015679	3.58	1.45	0.74	2.47	4.85			•			+
			•			(						-1

				Table 1-	4						
796754 AA460719	0.66	0.83	0.10	0.80	6.50					+ '	+
771053 AA430718	16.14		3.34	2.43	4.83						+
289071 N63609			7.78	2.45	4.82				.+		
637282 Al005515			0.70	1.55	5.71				+		
263697 H99672		0.93	1.58	4.57 .	2.68		+				
586780 AA130677		1.75	0.31	1.10	6.12				+		+
416374 W86202		2.66	0.81	1.69	5.52				+		
375682 AA032221		12.66	6.60	2.45	4.70						+
853968 AI791173		2.86	4.21	4.23	2.87		+				
287728 N75893		0.70	0.89	3.97	3.10	+					
B42927 AA486445		3.59	0.72	1.18	5.88				+	+	•
878281 AA670286	6.80	3.30	1.37	2.06	4.97					+	
359701 AA011176	5 5.21	2.32	1.09	2.25	4.78			•			+
897807 AA59853	5.08	2.77	0.98	1.83	5.16					+	
855749 AA663983	3 50.50	12.03	18.24	4.20	2.77			+			
264895 N2117	0 <b>7.86</b>	1.63	3.70	4.82	2.12			+			
825207 AA50412	0 11.96	2.92	4.19	4.09	2.86			+			
461476 AA70504	1 0.78	0.61	0.14	1.28	5.64		•			+	
449275 AA77770		0.83	1.64	4.58	2.32	+					
755239 AA42213		9.29	3.62	1.93	4.95						+
259973 N4782		1.95	0.28	0.85	5.99					+	+
290370 N6230		0.48	0.11	1.30	5.53						+
773287 AA42553	4 15.94	8.17	3.27	1.95	4.87						+
472689 AA87315	5.15	1.21	2.03	4.26	2.53	+	,				
252314 H8717	75 0.69	0.62	0.12	1.11	5.66						+
454317 AA67716	35 20.56	33.95	3.34	0.61	6.16				+		
1585549 AA9767	14 6.44	3.55	1.31	1.81	4.92						+
491764 AA1505	0.81	0.59	0.15	1.36	5.37						+
139771 R623	52 7.72	6.17	1.42	1.25	5.44			•	+		
462097 AA7053	76 8.32	6.33	1.55	1.31	5.38					+	
279482 N488	04 8.66	3.25	2.16	2.67	4.00				+		
824122 AA4906		2.75	0.94	1.69	4.95					+	
1031583 AA6093		9.06	15.59	4.20	2.44	+					
250869 N236		32.54	44.10	3.81	2.81	+				+	
896914 AA7793		7.40	4.35	2.44	4.14					т	+
814306 AA4591		10.68	3.77	1'.71	4.85	1					7
77915 T612		17.48	2.63	0.86	5.70	l			•		+
260116 N320		0.86	0.16	1.03	5.52			+			•
210405 H653		7.84	15.49	4.34	2.20			7			+
703943 AA278		2.81	0.99	1.70	4.82	1					+
824267 AA491		0.41	0.16	1.81	4.68	1		+			·
813973 AA455		0.88	1.18	3.70	2.75	1		•			+
825297 AA504			0.19	1.48	4.97	1					+
838807 AA457			0.33	0.74	5.71 4.14						+
1607765 AA989			0.96	2.29	3.58						+
	336 2.10		0.59	2.82					+		
1456160 AA862			7.76	1.03	5.36 2.12			+	•		
897767 AA598			0.98	4.25	4.63			•			+
1466631 AA883			0.26	1.71	5.28						+
	0.53		0.10	1.05		,			+		•
768643 AA430			9.10	1.74 0.98	4.57 5.32				•		+
610124 AA17			0.32	2.04	4.24						+
1631849 A1004			5.91 1.40	3.86	2.42			+			
204111 HS	5907 3.3	U.61	1.40	00.ر	2.42	- 1		-			

Page 4

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		_		Tabl	le 1-4						
491184 AA137144	6.35	1.83	2.27								
824237 AA491249	74.89	31.45	19.54	3.47	2.80	+					
24208 R38412	7.49	3.51	1.85	2.38	3.83						
267419 N24973	0.07	0.96	0.19	2.13	4.05						
731410 AA412250	0.53	0.58	0.10	1.01	5.16						+
869450 AA680244	51.58	35 <b>.87</b>	10.98	0.92	5.24						+
124252 R02439		4.74	1.25	1.44	4.70				+		+
593658 AA160080	1.08	1.02	0.21	1.28	4.85				•		
134969 R31789	6.81	4.59	1.48	1.07	5.04				•		
430465 AA680186	5.55	1.53	2.30	1.48	4.58				+		+
796351 AA459765	2.34	1.80	0.50	3.62	2.41	+			·		
549101 AA083485	12.81	2.43	17.02	1.30	4.72						
824376 AA489698	0.49	0.34	0.11	5.26	0.75		+	+			+
453599 AA679569	4.23	2.23	1.03	1.42	4.58			•		+	
530185 AA083671	8.46	7.83		1.89	4.10					•	
773248 AA425773	21.90	19.94	1.73	1.08	4.88					+	+
772416 AA405597	1.03	1.35	4.51	1.10	4.86					+	
1456721 AA864875	12.82	5.49	0.20	0.76	5.19					•	
50764 H17363	0.43	0.23	3.58	2.33	3.58						+
259462 N29545	3.69	1.65	0.11	1.89	4.00						+
366815 AA029490	71.61	18.49	1.02	2.24	3.62	-			+		+
731016 AA421256	0.85	1.00	36.97	3.87	1.94	+			•		
197779 R93515	8.61	6.61	0.17	0.85	4.96						
1519515 AA902831	0.85	0.55	1.91	1.30	4.50				+		+
511428 AA126009	24.31	24.09	0.20	1.55	4.25				•		
757206 AA443969	5.25	4.09	5.08	1.01	4.79						+
264465 N20522	0.77	0.84	1.17	1.28	4.47						+
609052 AA176607	0.87	0.84	0.16	0.91	4.79						+
1637343 AI015359	5.03	4.49	0.19	0.99	4.63						+
186768 H51914	9.43	5.75	1.12	1.12	4.50						+
613237 AA181767	1.07	0.78	2.39	1.64	3.95				+		+
841302 AA487213	7.41	3. <b>64</b>	0.25	1.37	4.22				•		_
251517 H96605	0.73	0.80	2.09	2.04	3.54				+		+
855624 AA664101	17.75	8.17	0.16	0.91	4.64				,		
841220 AA486723	1.46	1.23	5.31	2.17	3.34				+		+
1631194 AA994757	21.83	14.39	0.34	1.18	4.32				•		
251435 H97989	0.73	0.90	5.47	1.52	3.99						+
767406 AA417911	0.44	0.45	0.16	0.81	4.68					•	_
745283 AA625567	7.94	5.76	0.10	0.98	4.45						+
815017 AA465258	14.81	9.71	1.97	1.38	4.02						+
346889 W79834	0.89	0.21	3.84	1.52	3.85						+
823895 AA490490	1.57	1.72	0.79	4.21	1.12		+				+
1553306 AA934762	7.92	5.52	0.36 2.06	0.91	4.40						
361922 AA001403	9.18	5.49	2.55	1.43	3.84						•
290378 N64508	17.10	17.02	4.07	1.67	3.59						•
1602619 AA988798	135.32	117.37		1.00	4.20						
815800 AA485216	1.22	1.44	33.70	1.15	4.01						
377672 AA056001	2.59	1.39	0.29	0.84	4.25					+	
. 256680 н96392	1.72	1.47	0.81	1.87	3.20				+	7	
823578 AA497118	0.56	0.53	0.44	1.17	3.88				•		
878152 AA775431	4.20	3.49	0.14	1.06	3.98						
685922 AA262559	3.59	3.49 3.70	1.10	1.20	3.82						
786525 AA452113	1.46	0.42	0.90	0.97	3.99						
1594019 AA988701	24.27	24 <b>.53</b>	1.00 <b>6.</b> 55	3.46	1.46	+	,				
		~ 1.00	U.33	0.99	3.70						

				Table	: 1-4
282283 N52675	0.84	1.00	0.22	0.84	3.85
813271 AA455938	0.96	1.12	0.25	0.85	3.79
461336 AA699770	3.42	4.00	0.91	0.85	3.74
742581 AA400247	1.20	1.08	0.36	1.11	3.31
161456 H25546	13.35	10.78	5.08	1.24	2.63

Table 1-5

	GenBank	Aver.	<b>A</b>				
Clone	Ace No.	of	Aver.	Aver.	1	T/N	
282089		1.51	of BPH	0[	Spec Exp	& T/BPH	Secreted
646037		1.02	1.41	11.18	+	+	-
745283		5.76	0.75	24.42	+	+	4
30428		0.99	1.97	7.94	+		-
35058			1.16	4.29		+	+
46166		1.89	1.44	15.59		+	+
67401		0.93 0.20	1.71	13.13	,	+	+
240694		0.20	0.23	1.47		+	+
280217	N47941	5.86	0.83	4.53		+	+
292515	N68465	2.86	2.89	20.52		+	+
593185	AA159714	2.60 5.61	1.56	16.22		+	+
730739			2.29	28.06		+	+
1592006	AA953229	7.10	3.45	36.49		<b>`</b> +	+
32160	R43432	1.51	1.24	5.30		+	+
42415	R60981	3.59	2.77	26.80		+	
50188	H17943	0.51	0.72	6.86		+	
51773	H23211	2.29	2.53	10.75		, <b>+</b>	
81331	T60075	2.40	2.09	11.48		+	
222157	H85 <b>80</b> 6	0.78	0.51	2.84		+	•
281003	N50880	1.02	0.72	8.80		+	
281039	N47717	6.30	4.90	128.77		+	
283173	N45236	0.94	0.90	14.05		+	
308682	W25202	3.97	2.47	17.85		+	
450049	AA703396	0.87	0.27	8.99		+	
486035	AA043109	1.83	1.80	18.37		+	
488964	AA047260	3.47	1.86	24.75		+	
490178	AA121266	2.13	0.96	7.77		+	
712950	AA282272	5.63	2.73	24.55		+	
757435	AA437224	3.21	1.34	11.53		+	
784168	AA432103	59.86 7.37	58.35	365.78		+	
788180	AA453562		3.83	53.04		+	
814053	AA465495	0.81 1.70	0.50	25.18		+	
823940	AA490213	1.70	1.75	14.66		+	
868282	AA633957	2.01	2.83	22.39		+	•
877835	AA625634	4.34	1.47	8.99		+	
1034473	AA779728	1.51	3.34	23.06		+	
1034738	AA780190	3.90	1.35	33.61		+	
1048810	AA621342	3.04	3.28	22.40		+	
1412481	AA845156	3.50	2.59 0.90	20.15		+	
1556526	AA935560	0.15		8.41		+	
26519	R20669	19.25	0.12 3.76	0.88		+	
35626	R45292	1.09	5.76 6.51	36.05			+
51700	H22853	1.21	0.82	9.42			+
52933	H29315	1.99	0.58	3.66			+
77915	T61271	17.48	2.63	3.59			+
80050	T63893	3.84		14.99			+
84955	T74846	1.01	1.52	8.15			+
124252	R02439	4.74	1.43	5.26			+
134969	R31789.	4.59	1.25	6.06			+
		7.27	1.48	6.81			+

Table 1-5

_			_				
	GenBank	Ayer.	Aver.	Aver	Prostate	T/N	
Clone	Acc No.	of	of BPH	01	Spec Exp	& T/BPH	Secreted
150262	H01129	3.31	3.87	13.66			+
151055	H02230	7.51	4.62	27.32			+
159608	H16152	4.14	2.69	14.43			+
186768	H51914	5.75	2.39	9.43			+
186918	H43317	1.10	0.56	3.04			+ '
233419	H77706	1.14	1.90	5.55			+
251806	H96 <b>647</b>	1.83	0.46	2.94			+
252314	H87175	0.62	0.12	0.69			+
256680	H963 <b>92</b>	1.47	0:44	1.72			+
259462	N29545	1.65	1.02	3.69			+
260116	N32044	0.86	0.16	0.88			+
263697	H99672	0.93	1.58	4.23			+
280763	N50563	0.50	0.10	0.53			+
283312	N54763	6.63	12.14	63.94			+
284701	N64840	7.05	1.52	33.05		-	+
290370	N62301	0.48	0.11	0.62			+
290378	N64508	17.02	4.07	17.10		•	+
291255	N72215	1.35	4.65	15.20			+
293635	N63807	3.44	12.94	26.47			+
293651	N69653	1.84	6.56	13.35			+
321706	W35398	1.86	1.25	7.80			+
346889	W79834	0.21	0.79	0.89			+
361922	AA001403	5.49	2.55	9.18			+
375682	AA032221	12.66	6.60	31.04			+
377672	AA056001	1.39	0.81	2.59			+
416374	W862 <b>02</b>	2.66	0.81	4.49			+
430465	AA680186	1.53	2.30	5.55			+
450307	AA682851	1.62	2.25	9.39			+
454317	AA677165	33.95	3.34	20.56	}		+
461336	<b>AA</b> 699 <b>770</b>	4.00	0.91	3.42			.+
461476	AA705041	0.61	0.14	0.78	İ		+
491184	AA137144	1.83	2.27	6.35	<b>\</b>		+
491764	AA150502	0.59	0.15	0.81	i		+
509731	AA045699	23.94	6.63	39.41	l		+
509887	ľ	7.39	17.54	39.47	1		+
511428		24.09	5.08	24.31	l .		+
511952		92.92	45.17	225.66	}	•	+
530185	3	7.83		8.46	1		+
564801		3.96		12.35	i		+
586780		1.75		1.92	i		+
593658		1.02		1.08			+
594994		10.98		14.61			+
595695		2.33		3.58	2		+
609052		0.87		0.87			+
610124		1.73		1.70			+
666334		1.54		1.64			+
731410		0.58		0.53			+
756600		7.13		47.06			+
757392	AA426110	0.31	7 0.12	0.86	5 [		+

Table 1-5

ì	GenBank	Aver.	Aver.	Aver.	Prostate	T/N	
Clone	Acc No.	of	of BPH	of	Spec Exp	& T/BPH	Secreted
767406	AA417911	0.45	0.10	0.44			+
768643	AA430351	23.92	9.10	41.57			+
770066	AA430545	2.60	1.93	11.43			+
773248	AA425773	19.94	4.51	21.90			+
773479	AA427899	0.65	5.99	12.32			+
784130	AA432075	6.93	3 <i>.</i> 50	17.23			+
<b>787</b> 938	AA452278	1.27	1.08	5.09			+
788655	AA449837	2.78	1.58	7.66			+
<b>7952</b> 65	AA451807	1.26	0.97	4.58			+ .
796161	AA461092	5.27	15.52	30.31			+
796239	AA460675	23.22	4.18	26.07			+
811582	AA458516	5.34	1.75	10.13			+
811603	A1732784	2.74	6.22	19.62			+
811867	AA454963	2.97	7.43	24.57			+
815163	AA481144	0.57	0.19	1.20			+
823615	AA496957	3.54	1.97	9.40		-	+
823895	AA490490	1.72	0.36	1.57			+
824126	AA490611	0.37	0.10	0.59			+
825822	AA491370	9.48	1.97	14.07	i. I		+
826072	AA521394	0.83	0.18	1.20			+
826109	AA521327	0.54	0.10	0.75			+
838285	AA458747	1.36	0.21	1.34			+
838807	AA457681	2.58	0.33	1.90			+
839081	AA488732	5.37	2.78	16.60	:		+
840894 855624	AA482340	11.87	5.66	34.81			+
878152	AA664101 AA775431	8.17	5.31	17.75			+
884606		3.49	1.10	4.20			+
897767	_	13.42 0.49	5.43	33.46			+
951313	AA620591	6.68	0.98	2.07			+
1031583		9.06	1.01	6.84	1		+
1056172		4.78	15.59 1.77	38.01	l		+
1456160	1	40.32	7.76	10.33 41.61	l		+
1472689		1.21	2.03	5.15	1		+
1535596		285.67	66.91	401.67	ł		+
1610448		57.06	16.44	135.55	{		+
1631849		12.24	5.91	25.01	l		+
	1		3.71	23.01	1		+

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

AveTumor         Folid(AveTumor/Av Fold(MPM120IA Fold(MPM121IA Folid(MPM121IA Folid(MPM120IA) veBPH)         veBH)         veBH)         veBH)         veBH)	Table 2-1; T3NO vs BPH: Genes Upreguia tumor sample or 3 fold or higher in at leas	tumor sample or 3 fold or higher in at leas	Yord or night										
IMAGE   Colore   ID			3250			To the February For	MANA MANA E	old(MPM1211A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	veBpH) Samples	Samples
Clone ID         Number         133.47         67.99         149.63         165.70           788180         AA453310         1.88         113.47         67.89         149.63         146.70           646027         AA196878         2.38         114.00         47.88         662.4         24.62           646027         AA196878         2.38         114.00         47.88         66.24         24.62           245068         AA703386         2.55         7.64         30.08         38.52         11.57         4.60           245768         NGS80         2.40         4.23         17.34         20.64         4.13           245768         R35061         1.46         2.54         17.34         20.78         4.13           130608         R35061         1.48         25.46         17.34         20.78         4.13           130608         R43061         1.27         1.257         10.16         13.34         4.13           113600         R3508         A447747         1.24         1.24         11.34         4.13           113600         R3508         A451723         3.44         4.61         11.34         8.26           113600		IMAGE	Accession	AveBPH		Polo(AVE LUIIOIMA .	veBPH)	veBPH)	veBPH)	Value		•	•
788180         AAASS310         158         133.47         67.89         1420.0         47.83         66.24         2.62		Clone ID	Number				<b>\$</b>	185 70	7.38	52.32	22.54	10.36	<b>,</b> a
BACOTY         AA719687P         2.38         114,00         47.88         863.24         27.02           284701         N84840         2.41         73.10         30.37         31.27         60.07         34.24         47.88         863.24         50.07         42.05         42.05         78.49         30.08         38.52         11.57         42.05         42.36         17.34         22.19         11.57         20.04         47.38         17.34         22.19         11.57         20.04         47.34         17.34         22.19         11.57         20.04         47.34	٠	788180	AA453310	1.86	133.47	67.89	149.63	2,63	17.47	31.06	53.59	35.80	o <b>t</b>
264701         NABSADO         2.41         73.10         30.37         31.27         90.30           264702         AANDASSS         2.55         76.94         30.08         38.52         11.57         4.60           450048         AANDASSS         2.55         76.94         30.08         38.52         11.57         4.17           245068         AANDASSS         2.56         44.38         17.34         22.19         2.06         2.2           136060         R35051         2.56         44.38         17.34         22.19         2.064         2.2           136060         R35051         2.56         44.38         17.34         22.19         2.15         4.138         17.34         22.19         2.15         4.138         17.34         2.219         2.064         2.2         1.15         4.134	- 6	2001	44408070	2.38	114.00	47.98	65.24	70.67	24.87	6.28	41.24	19.50	۰ ه
450048         A.Z.03288         2.55         78.64         30.08         38.52         11.31         21.01           2.457048         N.52288         0.01         0.21         22.49         11.34         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         20.64         41.32         11.34         20.64         41.32         11.34         20.64         41.32         11.55         20.64         41.32         11.55         20.64         41.32         41.52         20.64         41.34         41.32         41.33         41.33         41.34         41.34	4 •	984704	N84840	2.41	73.10	30.37	31.27	30.02	48.94	16.67	45.42	20.45	<b>•</b>
245788         N.52389         0.01         0.21         22.49         11.34         8.60           245788         N.52389         2.05         423.65         17.62         22.19         20.64         2.06           281003         N.50880         2.05         423.65         17.23         20.48         41.32           130605         R.33061         2.66         44.38         17.34         20.48         41.32           133605         R.33061         2.66         44.38         17.23         20.48         41.34           133605         RA45723         3.44         46.10         11.39         11.55         50.28           119755         H51648         0.01         0.12         11.61         7.98         16.51           1102086         AA621761         1.24         12.57         10.16         7.98         16.51           1102086         NB1003         0.80         6.94         8.70         13.54         6.01           21831         17244         0.50         3.63         7.67         5.89         6.01           21831         17243         0.50         3.63         1.97         6.28         6.04           21831	, ,	180049	AAZOTABA	2,55	78.64	30.08	38.52	E 6	10.09	19.72	21.41	63.61	<b>5</b>
28/1000         N/50880         24.05         4.23.65         17.82         22.19         20.04           130606         R.35061         2.66         44.38         17.34         28.78         41.32           130606         R.35061         2.66         44.38         17.34         28.78         41.32           130606         R.35061         2.66         44.38         17.34         28.78         41.32           130708         R.28397         1.48         2.54.6         17.23         30.48         41.32           117975         H.5164         0.01         0.12         11.81         92.8         11.84           117975         H.5164         0.01         0.12         11.81         92.8         11.84           117076         1.57         10.16         7.99         15.78         15.14         15.14           15070         0.80         8.84         8.70         13.54         16.10           21831         172543         0.28         1.87         3.24         10.05           21831         172543         0.28         8.87         3.24         10.16           21831         172543         0.29         8.87         3.24	, ,	97476	NEGORB	0.0	0.21	22.49	11.34	20.00	22.40	15.57	11.61	13.29	φ.
136606         R35051         2.66         44.38         17.34         29.78         41.32           131600         R35051         2.66         44.38         17.23         30.48         41.34           131300         R28307         1.48         25.46         17.23         30.48         41.34           139808         AA457235         3.44         48.10         19.39         11.55         50.28           118753         H51649         0.01         0.12         11.81         92.8         11.84           118753         H51649         0.01         1.24         12.57         10.18         7.89         16.81           118703         0.80         6.94         8.70         6.78         6.78         6.03           118703         0.80         6.94         8.70         13.54         10.05         6.03           292217         N79180         0.80         6.94         8.70         5.89         6.00           292217         N79180         0.05         0.42         8.73         3.24         10.05           292217         N79180         0.06         6.03         3.87         3.84         4.64           292217         N79180 </th <th></th> <td></td> <td>NEGRED</td> <td>24.05</td> <td>423.65</td> <td>17.62</td> <td>22.19</td> <td>20.02 20.04</td> <td>0.28</td> <td>10.57</td> <td>5.71</td> <td>7.21</td> <td><b>1</b>0</td>			NEGRED	24.05	423.65	17.62	22.19	20.02 20.04	0.28	10.57	5.71	7.21	<b>1</b> 0
133130         RESISSY         148         25.46         17.23         30.48         41.54           838668         AA457235         3.44         46.10         13.39         11.55         50.28           179763         H51649         0.01         0.12         11.61         92.8         11.84           179763         H51649         0.01         0.12         11.61         7.89         16.61           105086         AA51781         1.24         12.57         10.16         6.78         6.28         11.84           1105086         AA61778         1.24         12.57         10.16         6.78         6.28         11.84           1105075         R81308         0.50         0.84         8.70         13.54         16.10         6.26         6.73         6.28         6.50         6.60         6.20 <th></th> <td>1863</td> <td>R35051</td> <td>2.58</td> <td>44.38</td> <td>17.34</td> <td>29.78</td> <td>70"14</td> <td>R 15</td> <td>9.24</td> <td>10.23</td> <td>5.98</td> <td><b>1</b>0 (</td>		1863	R35051	2.58	44.38	17.34	29.78	70"14	R 15	9.24	10.23	5.98	<b>1</b> 0 (
838688         AAA57235         3.44         46.10         13.39         11.55         90.00           179753         H51649         0.01         0.12         11.81         928         11.84           179753         H51649         0.01         0.12         11.81         928         11.84           170365         AA621761         1.24         12.57         10.16         7.89         16.61           51433         H20747         2.07         18.86         6.43         6.78         6.32           51433         H20747         2.07         18.66         8.70         13.54         16.10           306068         R91308         0.50         8.94         8.70         13.54         16.10           185075         R91308         0.50         3.83         7.67         5.89         6.60           22221         AN50484         1.91         12.18         6.34         6.03         6.54         8.44           22255         AA504484         1.91         12.18         6.03         6.34         4.64         3.19         6.01           22251         AA504484         1.91         12.18         6.03         6.54         8.46         4.64 <th>- «</th> <td>133130</td> <td>R28397</td> <td>1,48</td> <td>25.48</td> <td>17.23</td> <td>30.48</td> <td>45°C4</td> <td>58.</td> <td>4.31</td> <td>3.32</td> <td>7.01</td> <td><b>2</b>0 (</td>	- «	133130	R28397	1,48	25.48	17.23	30.48	45°C4	58.	4.31	3.32	7.01	<b>2</b> 0 (
179753         H\$1648         0.01         0.12         11,81         9.28         11,04           1030855         AA621781         1.24         12.57         10.16         7.89         16.81           51433         H\$1648         0.01         1.24         12.57         10.16         7.89         16.10           51433         H\$10747         2.07         19.56         6.94         8.70         6.78         6.32           306068         N91003         0.80         6.94         8.70         5.89         6.60           165075         R91398         0.50         3.83         7.07         5.89         6.60           21831         T7243         0.29         1.97         6.87         3.24         10.05           21831         T7243         0.29         0.42         6.73         3.34         10.11           292217         N79180         0.06         6.03         6.03         6.54         8.4           428816         AA60481         0.10         0.60         6.03         3.9         4.64           428816         AA60481         0.10         0.60         6.03         3.95         3.77         4.43           2	• •	BARRA	AA457235	8 4	48.10	13.39	11.55	07'00	11.99	10.01	17.09	10.66	<b>20</b> (
(CGOB65         AA621761         1.24         12.57         10.16         7.89         10.01           51433         H20747         2.07         19.56         9.43         6.78         6.32           306066         N91003         0.80         6.94         8.70         13.54         16.10           165075         R91308         0.50         3.83         7.67         5.89         6.01           165075         R91308         0.50         3.83         7.67         5.89         6.01           165075         R91308         0.50         1.97         6.73         6.29         6.01           222217         N79180         0.06         6.03         6.73         6.28         4.64           225277         N79180         0.06         6.03         6.37         6.28         4.64           225277         N79180         0.07         0.14         6.84         3.19         4.64           42816         AA004681         0.10         0.60         6.03         3.24         4.01           42816         AA004681         0.10         0.60         6.03         3.87         4.04           23071         AA01176         2.91	• \$	170753	HS1549	0.0	0.12	11.81	9.28	6	11.93	6.14	4.27	15.03	<b>20</b> (
51433         H20747         2.07         18.56         6.43         6.78         6.78         6.79           306066         N81003         0.80         6.94         8.70         13.54         16.10           165075         R91398         0.50         3.83         7.67         5.89         6.60           165075         R91398         0.50         3.63         7.67         5.89         16.10           21831         T72543         0.29         1.87         6.73         3.24         10.05           225217         N79180         0.08         0.42         6.73         6.28         4.64           825859         AA504784         1.91         12.19         6.37         6.28         4.64           428816         AA004681         0.10         0.60         6.03         6.34         4.64           428816         AA004681         0.10         0.60         6.03         3.19         4.64           428816         AA004681         0.10         0.60         6.03         3.21         3.84           428816         AA004681         0.10         0.60         6.03         3.21         3.87           428816         AA004681	2 \$	1020855	AA621781	1.24	12.57	10.16	7.99	19.01	65.4	7.12	8.55	18.24	<b>1</b> 0
306068         NB1003         0.80         6.94         8.70         13.54         10.10           195075         R91398         0.50         3.83         7.67         5.89         6.60           195075         R91398         0.50         3.83         7.67         5.89         6.60           21831         T72543         0.29         1.97         6.87         3.24         10.05           292217         N79180         0.08         0.42         6.73         3.94         10.11           292217         N79180         0.08         0.42         6.73         3.94         10.11           292217         N79180         0.08         0.42         6.73         3.94         10.11           292217         N79180         0.00         0.14         6.84         6.37         6.54         8.44           428616         AA004681         0.10         0.60         6.03         6.54         8.44           428617         0.02         0.14         6.84         3.19         6.01         4.08         3.17         4.13         3.87           24364         AA014176         2.91         12.87         4.42         3.17         4.13	= \$	61433	H20747	2.07	18.56	6.43	6.78	6.32	7.85	4.72	3.65	6.31	<b>6</b> 9 (
195975         R91398         0.50         3.83         7.67         5.89         8.60           21831         172543         0.29         1.87         6.87         3.24         10.05           21831         172543         0.29         1.87         6.87         3.24         10.05           292217         N79180         0.08         0.42         6.73         3.94         10.11           292217         N79180         0.08         0.42         6.37         6.28         4.64           42858         AA504784         1.91         12.18         6.37         6.54         8.44           428616         AA004681         0.10         0.60         6.03         6.54         8.44           42863         0.14         5.84         3.19         6.01         4.09         3.17         4.13           243641         AA014176         2.91         12.87         4.42         3.21         3.87           243641         AA014176         2.91         12.87         4.42         3.17         4.13           243641         AA014176         2.91         12.87         4.09         3.77         4.13           24364         AA32103	· •	306088	NB1003	0.80	6.94	8.70	13.54	16.10	8 8	5.51	12.71	6.41	<b>x</b>
21831         T72543         0.29         1.97         6.87         3.24         10.03           292217         N79180         0.08         0.42         6.73         3.94         10.11           292217         N79180         0.08         0.42         6.37         6.28         4.64           292517         N79180         0.08         0.42         6.37         6.28         4.64           428529         AA04484         1.91         12.19         6.34         3.19         6.01           428516         AA04681         0.10         0.60         6.03         6.54         8.44           428516         AA04681         0.10         0.60         6.03         6.34         6.01           428517         AA01178         2.91         12.87         4.02         3.17         5.98           243641         AA01178         2.91         12.87         4.02         3.17         4.13           243641         AA01178         2.91         12.87         4.06         3.17         4.13           243641         AA01178         2.91         12.87         4.06         3.77         4.13           24364         AA0203         15.97	2 \$	105075	R91398	0.50	3.83	7.67	5.89	9 6	08.6	7.18	5.75	8.20	<b>6</b> 0 1
292217         N79180         0.08         0.42         6.73         3.94         0.01           825858         AA504784         1.91         12.19         6.37         6.28         4.64           425858         AA504784         1.91         12.19         6.37         6.54         8.44           425816         AA604784         1.91         12.19         6.03         6.54         8.44           425816         AA04481         0.02         0.14         6.84         3.19         6.01           195034         R88764         0.02         0.14         6.84         3.19         6.01           24343         AA011776         2.91         12.87         4.42         3.21         3.87           2452036         H98688         0.22         0.86         3.95         3.77         4.13           2452036         H98688         0.22         0.86         3.95         3.77         4.13           2452036         H98688         0.30         3.88         12.94         16.20         2.45           2452036         H98688         0.30         3.88         12.94         16.70         2.45           258374         N31852         1.29 <th>. ÷</th> <td>21831</td> <td>172543</td> <td>0.29</td> <td>1.97</td> <td>6.87</td> <td>3.24</td> <td>0.01</td> <td>7.14</td> <td>10.04</td> <td>4.07</td> <td>5.11</td> <td><b>10</b></td>	. ÷	21831	172543	0.29	1.97	6.87	3.24	0.01	7.14	10.04	4.07	5.11	<b>10</b>
428559 AA504784 1.91 12.19 6.37 6.28 7.04 4.2 19.05 6.03 6.54 8.44 8.44 195034 R88764 0.02 0.14 6.84 3.19 6.01 3.19 6.01 3.87 3.87 4.2 3.21 3.87 3.87 4.2 3.21 3.87 4.2 3.21 3.87 4.13 2.2025 H88688 0.22 0.88 3.95 3.77 4.13 2.2025 H88688 0.22 0.88 12.94 15.20 2.45 12.84 AA32103 15.97 194.72 12.19 9.62 26.16 12.84 19.82	2 5	292217	N79180	90.0	0.42	6.73	3.94	7 7	5,38	5.13	5.23	11.57	<b>.</b>
428816 AA004681 0.10 0.60 6.03 6.54 0.11 195034 R88764 0.02 0.14 6.84 3.19 6.01 359701 AA011176 2.91 12.87 4.42 3.21 3.87 243641 M89548 1.55 0.34 4.09 3.17 5.98 243642 N82088 0.30 3.88 12.94 15.20 2.45 287528 N82088 0.30 3.88 13.54 10.47 7.91 24.39 210820 H87712 0.19 1.82 8.61 2.75 9.06 210820 H87712 0.09 0.74 8.39 8.46 28.73 210850 AA809485 0.27 2.07 7.85 5.89 10.86 288211 NZ8988 0.57 3.71 6.48 8.59 3.89	2 定	825859	AA504784	1,91	12.19	6.37	6.28	5 6	9.60	4.79	4.07	3.73	<b>.</b>
195034 R88764 0.02 0.14 5.84 3.19 0.01 359701 AA011176 2.91 12.87 4.42 3.21 3.67 243547 M49548 1.55 0.34 4.09 3.17 5.98 243547 M49548 0.32 0.88 3.95 3.77 4.13 252036 H98698 0.30 3.89 12.84 16.20 2.45 287528 N82088 0.30 3.89 12.84 16.20 2.45 287528 N82088 0.30 15.97 164.72 12.19 8.62 255374 N31952 1.29 13.52 10.47 7.91 24.39 255374 N31952 1.29 13.52 10.47 7.91 24.39 259374 N31952 0.19 1.82 8.61 2.75 8.06 1 210320 H87712 0.19 1.82 8.61 2.75 8.06 255374 N31952 0.00 0.74 8.39 8.46 28.73 256311 N29988 0.57 3.71 11.25 6.59 7.39 3.89 266211 N29988 0.57 3.71 6.48 8.59 3.89	: 5	428816	AA004881	0.10	0.60	6.03	45.5	‡	2	7.48	3.67	8.48	<b>.</b>
358701 AA011178 2.91 12.87 4.42 3.41 5.98 2436.41 N495.48 1.55 6.34 4.09 3.17 5.98 282036 H96686 0.22 0.88 3.95 3.77 4.13 282036 H96686 0.30 3.89 12.94 16.20 2.45 287528 N82096 0.30 3.89 12.94 16.20 2.45 287528 N82096 0.30 3.89 12.94 16.20 2.45 259374 N31952 1.29 13.52 10.47 7.91 24.39 259374 N31952 1.29 13.52 10.47 7.91 2.75 9.06 259374 N31952 0.09 0.74 8.39 8.46 26.73 269160 772336 1.71 11.26 6.59 7.39 3.89 268211 N29886 0.57 3.71 6.48 8.59 3.89	2 2	195034	R88764	0.02	0.14	5.84	3.19	, c	3.89	8.86	3.53	3.08	<b>.</b>
243841         NA9548         1.55         6.34         4.08         3.17         4.13           262036         H96686         0.22         0.88         3.95         3.77         4.13           287528         N62066         0.30         3.88         12.94         15.20         2.45           287528         N62066         0.30         3.88         12.94         15.20         2.45           25873         AA432103         15.97         194.72         12.19         8.62         26.16           210820         H67712         0.19         1.35         10.47         7.91         24.39           210820         H67712         0.19         1.82         8.61         2.75         9.06           193923         R83407         17.59         158.73         8.02         10.87         16.00           1031599         AA609485         0.27         2.07         7.65         5.89         10.96           266211         N29986         0.57         3.71         6.48         8.59         3.89	8	359701	AA011178	2.91	12.87	4.42	3.21	, E	4.16	3.84	3.52	3.85	p 9
262035         H98686         0.22         0.86         3.95         3.17           287528         N62086         0.30         3.89         12.94         16.20         245           287528         N62086         0.30         3.89         12.94         16.20         245           259374         N31952         1.29         13.52         10.47         7.91         24.39           210820         H67712         0.18         1.82         8.61         2.75         9.06           193823         R63407         17.59         158.73         8.02         10.87         18.00           193829         AAB09485         0.27         2.07         7.85         5.80         10.96           103159         AAB09485         0.27         2.07         7.85         5.80         10.96           266211         N29986         0.57         3.71         6.48         8.59         3.89	2	243841	N49548	1.55	45.0	4.08	; ; ;	4.13	3,55	4,47	3.77	5	o <b>v</b>
287528         N82086         0.30         3.88         12.94         12.54         15.57           784168         AA432103         15.87         194.72         12.19         9.62         26.16           258374         N31952         1.29         13.52         10.47         7.81         24.39           210820         H67712         0.19         1.82         8.61         2.75         9.06           210820         H67712         0.19         1.82         8.61         2.75         9.06           193823         R83407         17.59         158.73         9.02         10.87         18.00           1931539         AA809485         0.27         2.07         7.65         5.80         10.86           1031589         AA809485         0.27         2.07         7.65         5.80         10.86           28610         172336         1.71         11.26         6.69         7.39         5.16           286211         N29986         0.57         3.71         6.48         8.59         3.86	ឧ	262036	H98668	0.22	0.88	55 E	3.7.	2.45	18.84	12.53	18.48	5.6	<b>.</b>
784168         AA432103         15.07         194.72         12.19         12.13         24.39           258374         N31652         1.29         13.52         10.47         7.91         24.39           210820         H67712         0.19         1.82         8.61         2.75         9.06           183823         R83407         17.59         158.73         9.02         10.87         18.00           183820         R83407         17.59         158.73         9.02         9.46         2.8.73           37830         R81372         0.09         0.74         8.39         9.46         2.8.73           1031598         AA608485         0.27         2.07         7.85         5.69         10.86           28610         17230         1.71         11.26         8.59         7.39         5.16           286211         N2986         0.57         3.71         6.48         8.59         3.86	ឌ	287528	N62098	0.30	3.88	12.94	16.20	26.16	14.54	17.70	<u>4</u>	3.92	n 4
258374         N31852         1.29         13.52         10.47         1.29         606           210820         H87712         0.16         1.82         8.61         2.75         9.06           183823         R83407         17.59         158.73         8.02         10.87         18.00           37980         R81372         0.09         0.74         8.39         9.46         28.73           1031599         AAB08485         0.27         2.07         7.65         5.89         10.86           26160         T72336         1.71         11.26         6.59         7.39         5.16           266211         N2988         0.57         3.71         6.48         8.58         3.86	75	784168	AA432103	15.97	194.72	12.19	7 01	24.39	10.24	5.27	13.26	1.75	5 K
210820         H87712         0.18         1.82         9.61         2.73         18.00           193823         R83407         17.59         158,73         8.02         10.87         18.00           37880         R81372         0.09         0.74         8.39         8.46         28.73           1031589         AAB09485         0.27         2.07         7.65         5.89         10.86           28180         T72336         1.71         11.25         6.59         7.39         5.16           286211         N29886         0.57         3.71         6.49         8.59         3.86	25	259374	N31852	1.29	13.52	10.47		9.08	3.94	3.08	33.98	6.6	, w
193823 R83407 17.59 158.73 8.04 10.05 28.73 37880 R81372 0.09 0.74 8.39 9.46 2.8.73 10.36 10.31599 AA809485 0.27 2.07 7.85 5.89 7.39 5.16 86180 T72336 1.71 11.25 6.59 7.39 3.86 28211 N29886 0.57 3.71 6.48 8.58	58	210820	HB7712	0.19	1.82	19:8	£. £7	18.00	7.67	13.10	22	3.2/	, <b>k</b>
37980 R81372 0.09 0.74 8.39 5.59 10.96 1031599 AA809485 0.27 2.07 7.65 5.69 7.39 5.16 89180 T72336 1.71 11.26 6.59 7.39 8.59 3.86 288211 N29888 0.57 3.71 6.48 8.59	27	193923	R83407	17.59	158.73	20.8	0.0	28.73	3.79	3.02	4.81	t ()	s w
1031599 AAB09485 0.27 2.07 (.85 7.39 86160 T72336 1.71 11.26 6.59 7.39 288211 NZ9986 0.57 3.71 6.48 8.59	8	37880	R81372	0.00	0.74	8.38	 	10.96	6.72	12.64	2.83	80.	о w
86160 T72336 1,71 11,25 0,58 1,55 288211 N29886 0,57 3,71 6,48 8,59	2	1031599	•	0.27	2.07	69.7	23.3	5.18	8.	15.83	2.68		. v
266211 N28986 0.57 3.71 0.40	ຊ	86160	172338	1.7	11.26	80.0		3.88	13.77	2.12	8.3	5	,
	E	268211	N29986	0.57	3.7	ġ.	2	•					

Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	GE Gen Bank BID Accession Number	Avebph	AveTumor	Fold(AveTumorian Fold(MPM120/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # of Tume • BPPH) VARPHI VERPHI CERTIC	Fold(MPM120/A I	Fold(MPM121/A )	Fold(MPM124/A	*old(MPM122/A	Fold(MPM123/A	Fold(MPM125/A	# of Tumc
•	32 48748R			1	•	· ·		(uuan	VeBPH)	VeBPH)	VeBPH)	Samples
- •			0.38	2.39	<b>7</b> .0	5.85	10.82	5.48	10.02	ş	,	•
	_		3.28	20.49	6.28	1.91	3.57	4.27	14.38	790	2.00	- o . :
•		EZ AA454587	3.59	21.22	5.82	9.12	7.48	7		5 6	<b>T</b>	•0
43)	35 428436	36 AA004412	0.53	3.12	5.83		3 4	Ç !	3.0	2.69	8.45	40
60	36 130138	36 R20813	9.0	0.20	5.83	2 2	) ) ; ;	4.73	5.71	11.11	7.68	45
•	37 347586		6.0	0.24	0.02 6 83	S7 /	08.60	5.30	5.38	6.96	2.08	w
8	8 785729	1	6	***	70.0	5A.2	4.97	4.10	9.4¥	3.72	11.48	9
88			7 .		86.6	6.69	4.98	0.83	11.93	8.34	3.73	40
<b>Ş</b>		-	00.00	(C.5)	5.51	3.24	6.98	2.95	6.31	8.01	4.56	<b>•</b>
4			3 8	20.	5.39	2.31	3.63	6.60	3.61	13.09	3.10	, <sub>10</sub>
. 3			0.83	4.38	530	2.79	3.52	3.68	421	80.6	75	) <b>v</b>
: 5		•	£7.7	9.09	5.25	3.31	10.29	3.87	2.38	2.58		, •
			8.	5.43	5.10	3.95	6.61	6.92	808	278	) ;	<b>.</b>
; 5			1.68	8.62	5.08	4.26	6.83	4.57	7.80	4.62	5 5	o <b>u</b>
? 9	_	_	Kį.	6.16	4.82	2.53	5.09	3.89	4.48	57 B	2 3	D 1
? !			¥.7	23.13	4.92	2.33	3.91	3.68	11 43	? ?	~ (i	<b>o</b>
<b>;</b> ;	134718		0.28	1.40	4.85	5.02	7.58	807	2 4	7.05	D7.4	kr) i
*	287745		3.81	18.41	4.83	2.38	4.28	375		2 6	2.83	<b>~</b>
4	271141	N34482	0.01	0.06	4.58	0.78			\$ 1	3.82	5.12	<b>\$</b>
8	1240414	4 AA788780	0.79	3.60	4 55		0.00	4.18	3.78	5.37	4.58	•
2	307249		338	15.28	3	90. L	4.13	3.48	7.19	6.3 <u>4</u>	4.47	40
. 25	783849	•	0.13	0.50	\$ 9	872	5.31	0.40	5.39	3.29	3.63	50
8	807720	AA598982	0.32	4. 5. 2. 2. 2. 2.	8.48	4.82	5.19	6.45	4.87	1.71	2.	wo
3	283070			90.1	4°7	3.68	3.50	2.84	3.83	6.48	5.83	¥
8	285005		3 6	0.12	4.31	3.87	6.96	3.48	6.69	154	4,45	<b>•</b>
8	343887		20.00	02.0	4.23	6.18	3.28	3.22	6.00	0.56	6.16	, NO
6	416855			)	4.16	6.16	7.98	3.20	3.85	0.24	3.50	
88	842848	_	, t	50:1	£.03	2.15	7.06	3.45	3.09	6.14	331	• •0
20	40689			<b>3</b> 0	<b>4.02</b>	3,49	4.63	2.84	***	5.85	3.05	· •:
8	358681	_	_	07.00	3.70	3.25	3.93	4.50	3.68	4.50	2.31	
5	128230			70°C	3.64	4.70	6.58	3.16	3.29	3.49	0.64	• •
B	375682		_		3.50	3.18	3.36	3.88	4.60	4.89	0.89	• •c
8	1488280		ь.	45.43	2.80	3.03	3.29	3.61	3.03	3.10	1.46	
2	741070		_	22.38	32.16	18.89	1.69	59.34	105.24	0.80	£ 61 00 61	•
<b>*</b>	130046			12.90	22.20	29.78	48.11			23.62	4.26	
;	5	AA643582 1.		33.09	18.02	49.03	18.43	32	•	2.77	3 5	• •
						r						•

Table 2-1;T3NO vs. BPH; Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			FELLIAT	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	Samples
,	MAGE	Accession	AveBPH	Ave Tumor	rotateve i untolitore •BPH)	veBPH)	veBPH)	VeBPH)	veBPH)	VeBPH)		
		Number			•		10 0	7	57.03	15.40	1.72	₹
8	41647	R52794	0.40	6.3 24	15.84	8.65	- 0.5	. 4	8.52	48.21	5.35	•
6	7484	AA625890	1.90	22.42	11.81	2.76	 	200.	8.26	2.32	1.91	4
88	1034473	AA779728	0.13	1.38	10.50	22.49	23.74	97.6	13.87	1.00	1.00	4
8	487165	AA045074	0.0	0.09	8.98	13.49	10.45	6.6	8.33	2,	2.78	4
2	1458424	AA862992	0.08	0.69	8.20	10.16	21.53	6. c	24.39	4.57	6.15	۳
	360155	AA013260	0.01	0.11	8.02	10.07	0.73	02.2	10.48	19:	2.63	4
2	144740	R78229	0.19	1.27	6.55	7.58	10.38	0.02	2.85	10.25	15.67	4
ß	24915	R39066	0.93	5.97	6.42	3.07	2.31	) ; ; ;	08.6	2.35	3.29	4
7	593185	AA165678	. 02.8	60.24	6.21	9.37	12.99	2.40	, c	1.74	10.42	4
73	1056172	AA620995	11.61	70.00	6.03	8.66	5.08	C. 2	5.02	1,51	2.67	•
78	782383	AA431407	0.95	5.55	5.84	8.95	41.77		7000	2.05	3.18	4
: =	1031113	AA609914	0.01	0.07	5.65	16.58	0.83	2 2	10.4	5.07	2.77	4
. 82	461541	AA705258	0.3	1.88	5.48	2.60	6.47		? C	2.29	2.67	*
67	491708	AA115259	0.75	3.90	5.23	3.93	6.17	F., 7	4.30	0.26	5.98	. ▼
2 2	787938	AA452278	1.76	8.87	5.04	6.20	9.84	6.65	) (2 (4 (4)	79.9	9.17	4
: <b>=</b>	754034	AA479058	0.05	0.10	5.01	5.74	6.67		70.0	1.1	3.26	*
. 28	970580	AA883077	9.	0.18	4.93	3.02	13.78	4.82	6 80	0.64	<b>4</b> .09	4
2	68787	153389	0.02	90.0	4.91	1.66	3.59	10.99	9.36		2.87	4
3	292515	N68465	3.53	17.34	4.91	3.22	3.1	2.03	11.08 0.11	2.70	10.45	4
<b>.</b> 2	298210	N74392	0. 4	2.17	₹.80	3.19	3.94	6.56	7. F	1.57	11.65	4
2	40402	R52089	0.02	0.09	4.82	3.44	0.53	59.4	71.1	4.42	5.71	4
. 6	858153	AA633811	0.05	0.24	4.82	1.99	4.80	Z.74	07.6	90%	123	4
5 8	704897	AA282253	0.27	1.26	4.71	1.92	9.73	7.83	3.55	 	207	4
2	814308	AA459318	9.65	45.42	4.71	5.23	10.10	19.7	07.0	1.30	7.69	4
8	781233	AA448318	0.02	0.10	4.70	0.48	<b>4</b> .83	0.0 0.0		1.85	2.40	4
2	823794	AA490283	1.70	<b>2</b> .	4.66	3.80	10.81	9. 69. 69. 69.	200	1.63	₩.	4
8	280170	N32072	12.85	58.29	4.61	5.52	5.50	3. 5		2.38	2.87	4
8	449309	AA7777802	2.47	11.22	4.54	4.85	3.65	3.67	1.78	184	3.78	4
Z	236155	H81758	0.98	4.35	4.51	<del>4</del> 2	2.42	10.20	07.9	9.60	3.44	•
68	296679	W02258	9.9	4.15	4:41	1.85	241	2 6	1.35	5.02	1.7	•
8	195387	R88915	0.47	2.07	4.38	¥.	G.46	9. c	1.35	3.86	6.85	~
26	428267	AA004628	1.41	6.20	4.38	2.17	5.43		10.05	2.98	1.97	•
88	391984	A1003825	4.42	19.15	4.34	3.25	4	, t	9.71	1.81	3. 3.	•
8	61700	H22854	1.82	8.25	4.29	5.18 61	<b>4</b> .3/	2	;			

	MAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumot/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor ebpt) veBpt) veBpt) veBpt) veBpt) veBpt) veBpt) veBpt) veBpt)	Fold(MPM120/A	Fold(MPM121/A veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A 1 vebpH)	Fold(MPM126/A veBPH)	Samples
;		Number	;	, ,		a V	68	2.50	3.26	4.67	2.90	
\$	754284	AA479484	12.01	62.10	<b>V</b> . :	0 10		02	7.43	8,18	1.28	•
<del>5</del>	210697	H66840	0.75	3.08		1.97	0.85	 	2.50	1,75	4,48	~
42	809456	AA443080	0.02	0.10	01.4	200	80 6	3.63	5	4.28	2.07	•
\$	361659	W96189	6.37	8 ;	B C	9 6	248	4.08	2.17	8.08	3.5	4
ጅ :	148225	H13688	27.5	,0; ;	4.07	50.00 00.00 00.00	3.21	2.71	10.52	69.0	3.20	4
5	417867	W80128	13.56	8.72	<u> </u>	9.00	5 6	3.88	5.13	8,53	2.31	4
\$ ;	436531	AA703048	<u>.</u>	5.81		1.41		898	2.74	2.97	3.41	•
<u> </u>	1031348	AA609106	<b>*</b>	<b>4</b> 6	2 C			11.7	3.69	6.17	1.75	•
<b>3</b>	277777	A4939275	5.0	00°7	- 0	20.4 20.6	, e,	<u> </u>	1.37	4,41	3,7	<b>~</b>
2 5	144/4/	K/624/	? ?	)	9 c	2, c	47.4	3.81	2.10	3.38	2.84	•
2 ;	301300	WSZ472	3 5	÷ <	3.18	2.57	4.47	6.19	3.29	7.01	0.18	4
= {	20707	AACU142	3 6	<u>.</u> .	0.78 87 E	6.40	3,53	3.79	6.85	1.28	0.83	•
2 5	40407 088800	C00C7N	3 5	÷ 4	9.76	2.83	4.18	2.58	3.38	4.60	<b>3</b> .	<b>~</b>
? ?	283005	NEGGTR	880	3.42	366	2.34	3.33	3.62	2.44	4.10	5.85	<b>4</b>
¥	350784	AA011320	3.58	13.08	3.65	4.67	5.61	2.55	4.74	3.85	0.48	4
: =	854701	94830104	0.88	3.58	3.65	2.92	5.15	3.36	5.17	4.30	9.	4
: ‡	24888	HR2081	2 E	3.03	3.63	4.16	2.78	3.62	5.06	3.32	2.83	•
. 4	77,1053	AA427528	10.52	38.08	3.62	3,55	5.47	3.81	5.38	2.39	=	4
5 5	488010	AA045709	0.98	35.5	3.60	1.73	6.29	3.11	1.20	3.29	5.98	•
120	112482	TROBBO	11.33	40.58	3.58	1.21	4.61	5.01	4.80	4.57	1.28	₹
2	88035	TR2842	0.03	0.10	3.58	5.81	3.02	3.15	2.19	2.78	<b>₹</b>	<b>~</b>
122	855832	AA664104	0.86	3.38	3.52	2.09	3.77	3.88	1.82	5.82	3.66	₹
<u> </u>	811914	AA454988	2.96	10.39	3.51	3.92	5.10	3.87	3.79	2.79	1.62	•
<b>1</b>	281371	NATABA	0.36	125	3.49	5.39	3.04	3.53	2.17	2.42	<del>,</del>	*
2	383.623	AA704850	9.01	31.38	3.48	2.44	3.12	4.85	1.73	3.48	620	₩
2	395400	AA757404	2.20	7.86	3.48	2.26	5.31	3.00	4.08	4.15	2.08	*
127	138507	R63253	14.	88,	3.47	1.74	4.81	3.90	3.23	4,16.	2.85	<b>~</b>
128	289071	N83609	19.65	68.00	3.46	3.37	5.09	1.38	1.59	6.09	3.28	₩.
8	480203	AA877461	0.55	1.89	3.45	1.61	4.23	4.13	4.28	4.83	1.55	▼ ·
8	242081	H83335	2.71	9.21	3.40	5.12	6.73	3,43	3.25	0.82	0.83	•
5	207098	H48502	0.16	870	3.38	3.02	4.77	3.04	2.28	4.65	752	₹ -
132	396839	AA758154	0.68	2.24	3.36	1.87	5.62	3.24	4.60	1.60	3.25	•
\$	209381	H84095	0.97	3.25	3.35	2.04	3.28	4.07	1.86	€.80	3.88	₹

•	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor ABPH) veBPH) veBPH) veBPH) veBPH) veBPH)	Fold(MPM120/A veBPH)	Fold(MPM121/A F	old(MPM124/A	Fold(MPM122/A veBPH)	Fold(MPM1ZS/A   veBPH)	Fold(MPM126/A veBPH)	Samples
	Clove ID	Number						88	3.28	1.47	3.18	· <b>→</b>
2	23807	R38364	0.38	1.24	3.28	7.98 86.1	- A.O.	3 4	3.37	3.87	2.04	*
\$5	1407905	AA883504	1.18	3.84	3.27	1.87	3.82		68.0	4.48	1,12	₹
=======================================	1858528	AA935560	1.17	3.77	3.21	3.58	5.83	0.40	¥38	3.33	3.88	•
<b>†</b>	491418	AA115537	9.54	1.72	3.20	1.82	3.32	0 6	4.18	5.11	1.80	4
\$	744001	AA829039	3.87	12.24	3.17	1.60	3.03	77.	3.86	3.55	1.84	▼
3 5	293110	N63848	3.38	10.70	3.17	2.38	3,94	<b>4</b>	2.89	3.22	3.21	•
\$ \$	782047	AA429602	1,17	3.67	3.14	3.26	2.51	3.73	3.87	3.10	4.58	4
<u> </u>	454689	AA877206	1.02	3.16	3.08	2.09	3.15	80°1	3.20	1.06	1.87	<b>~</b>
142	384293	AA709370	0.50	£.	3.05	4.26	, i	. c	3.14	3.71	1.62	4
3	34081	R44132	0.63	1.63	3.04	2.04	3.94	. c	4.18	0.88	4.58	*
3	743382	AA400385	0.03	0.09	3.02	3.33	3.10	, co	3.17	3.53	3.40	•
<u> </u>	128785	R10015	4.1	3.4	3.02	2.11	2.81	3.08		2.22	3.84	•
3 4	429298	AA007370	1.28	3.83	2.98	3.22	3.29	20.C	3.12	1.85	3.33	4
7	136818	R36465	0.47	1.37	2.94	3.15	2.85	3.25	1.72	0.47	3.55	4
3	433350	AA700604	42.83	123.16	2.88	3.66	5.22	3.75	<u> </u>	0.38	3.03	•
9	755276	AA496349	0.11	0:30	2.74	3.57	4,62	3.03	14.34	134,44	1,13	ຕ
3	270917	N32514	0.01	0.23	25.82	3.34	1,13	2 7	0.61	3.63	22.02	63
55	782233	AA431721	0.20	3.10	15.42	64.92	1.22	- 60	71.38	3.78	2.68	က
152	320495	W16659	0.08	0.88	13.99	1.39	3.75	7 2	28.47	44.95	2.05	က
153	293578	N94111	0.96	12.90	13.42	0.73	1.52	; ; ;	2.22	0.30	1.22	m
3	281038	71774N	0.78	8.89	12.85	43.16	25.22	- F	0.27	18.31	0.82	က
155	490718	AA115761	2	0.48	12.37	<b>46.34</b>	6.67		55.00	5.81	ы 1	ю
\$5	796747	AA443147	0.08	0.82	11.12	0.12	21.0	7.51	1.25	0.88	1.67	6
157	287728	N59158	0.51	5.43	10.68	20.89	30.40	15.4	1.88	1.07	2.28	n
\$	282089	N48259	1.90	19.34	10,17	Z1.69	5 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	18	1.28	0.70	<del>*</del>	m
159	731290	AA418827	4.08	33.97	8.32	14.84	4.75	1.05	5.39	32.28	2.68	<b>6</b>
\$	470216	AA028997	2.95	23.82	8.02	/6		163	1.38	34.08	2.42	0
5	375827	AA039851	0.07	0.57	7.75	K	3.42	1.15	32.30	5.68	0.12	n
162	277714	N48577	0.13	<b>7</b> 6.0	7,33	0.87	2,03	1 10	3.88	0.99	1.36	n
5	757451	AA437241	0.47	3.37	7.13	4.67	20.78	2.60	21.59	8.58	1.59	က
\$	488035	AA040881	5.25	35.89	6.84	2.41	45.62	22	1.58	1.47	3.24	6
165	299417	N74131	0.08	0.48	6.33	13.15	20.02 80.02	4.25	20.28	2.28	0.81	က
\$	666218	AA233809	9.6	0.23	8.24	1.91	Da: 1	9 0	4.75	0.58	22.68	က
187	757389	A4437142	0.02	0.11	5.78	1.70		3				
							s.					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor semi----

144         185         184         185         440         734         734           149         180783         182352         0.83         1.43         1.43         1.47         1.52         4.09         734           149         172284         48242277         0.83         1.45         5.89         6.89         1.47         1.72         0.03         0.73         0.03         0.73         0.03         0.73         0.03         0.73         0.03         0.73         0.03<		INAGE Clone ID	Gen Bank Accession Number	AveBPH	AveTumor	Fold(AveTumor/Av Fold(MPM121/A Fold(MPM121/A Fold(MPM122/A Fold(MPM121/A Fold(MPM126/A \$ of Tum	Fold(MPM120/A veBPH)	Fold(MPM1211A   veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A vebPH)	Fold(MPM126/ veBPH)	Sample:
90 CATTABOL ALCASSATY         0.81         4.58         6.88         1.65         4.47         1.15         1.10           178 DAGA ALCASTATA         5.80         0.22         0.28         6.58         6.81         1.42         8.82         0.17         0.10           2 224.63         0.22         1.28         6.59         6.43         6.43         0.43         1.77         0.13         1.10           2 224.64         VH6424         0.22         1.23         6.50         0.43         4.63         0.73         0.03         0.48           7 72537         AA444464         1.72         0.40         4.73         7.20         0.20         0.48         1.14         0.09         1.14           7 72537         AA442660         8.71         4.00         4.83         1.82         1.24         0.73         1.14         1.14           8 77527         AA442660         8.71         4.00         4.83         1.82         1.80         0.93         1.43         1.80         0.93         1.14         0.93         1.14         0.93         1.14         0.93         1.14         0.93         1.14         0.93         1.14         0.93         0.93         0.93 <th< th=""><th>168</th><th>198168</th><th>R92362</th><th>0.59</th><th>3.32</th><th>5.80</th><th>1.71</th><th>1.65</th><th>2.16</th><th>16.65</th><th>4.09</th><th>7.3</th><th>m</th></th<>	168	198168	R92362	0.59	3.32	5.80	1.71	1.65	2.16	16.65	4.09	7.3	m
0         712850         AACASCT73         5.93         3.2.98         6.84         14.24         8.82         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.74         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.78         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.77         0.49         0.78	169	687667	AA235370	0.81	4.56	5.59	5.88	18.59	4.47	1.32	1.05	2.26	က
1         500781         AAA448778         0.08         0.43         5.46         0.051         7772         0.13         1797         1797           2         2.24243         Nig624         0.22         1.23         6.30         2.13         6.09         0.73         2.09         0.73         1.18         0.09         0.13         1.14         1.18         1.14         1.24         1.25         6.09         2.73         6.09         0.73         0.09         0.74         1.18         0.74         0.09         1.13         1.14         1.24         1.25         0.09         0.74         1.14         1.24         1.24         1.25         0.09         0.14	170	712950	AA282273	5.83	32.98	5.58	6.81	14.24	8.62	0.72	9870	2.12	က
2 323445         WH6424         0.23         1.23         6.30         3.11         4.09         0.073         20.03         0.148           2 3 3501         RA41164         1.22         6.30         5.11         6.30         2.72         4.45         2.60         1.48         7.48           2 77254         RA441644         1.12         6.00         5.01         6.01         4.49         7.24         4.97         7.68         1.21         2.77         0.09         1.13           7 73254         RA422660         8.71         4.00         4.49         1.62         1.21         2.77         0.09         1.13           7 340665         AA422660         8.71         4.70         4.62         1.62         1.21         2.77         0.09         1.13           7 340665         AA422660         8.71         4.62         1.62         1.62         1.61         0.69         1.72         1.73         1.73         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83         1.74         1.83	171	504781	AA148735	0.08	0.43	5.45	6.26	0.51	7.72	0.13	17.97	0.10	က
4         735571         A4446444         1.72         4.85         2.00         1.65         11.48           4         725557         A4446444         1.72         4.97	Ē	322443	W18424	0.23	1.23	6.30	3.13	4.09	0.73	20.36	0.89	2.62	က
4 772554         AAAA46044         1172         8.00         6.01         2.62         2.75         8.00         0.89         7.44           6 777256         NB7002         1.48         7.34         4.07         7.69         11.21         2.17         0.90         1.13           8 77226         NAC2569         8.47         1.09         4.68         1.82         1.20         6.64         1.01         0.09         1.13           8 24756         AAA2569         8.47         0.04         2.09         4.64         1.68         1.42         1.91         4.83         0.26           8 24757         AA3264         0.04         0.05         4.46         1.68         1.47         1.93         0.26         0.26           7 2585         AA156         0.06         0.03         4.52         0.07         1.93         0.04         1.93         0.04           7 2586         AA156         1.67         1.69         1.47         1.49         1.83         0.04         1.93         0.04           7 2586         AA156         1.67         1.68         1.47         1.43         1.83         0.04         0.04         1.83         0.04         1.93	Ē	35010	R45118	0.66	3.31	5.03	2.72	4.85	2.60	1.65	11.48	8.86	60
6         277200         NATZGAGO         1.48         7.34         4.87         7.88         12.12         2.17         0.00         1.13           2 478208         AAAZGAGO         8.71         4.70         4.85         1.82         1.80         6.64         12.13         2.88           2 47820         AAAZGAGO         8.71         4.70         4.85         1.82         1.80         6.64         1.21         2.88         1.88         2.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.89         1.89         6.64         1.89         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.49         1.89         6.69         6.07         1.89         1.89         6.69         8.03         6.74         1.89         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.88         1.89         6.69         8.03         6.74         1.89         1.89         1.89         1.89         1.89         1.89 <th>7.</th> <th>782537</th> <th>AA448484</th> <th>1.72</th> <th>9:00</th> <th>5.01</th> <th>2.52</th> <th>2.75</th> <th>8.08</th> <th>0.99</th> <th>7.48</th> <th>8.28</th> <th>e</th>	7.	782537	AA448484	1.72	9:00	5.01	2.52	2.75	8.08	0.99	7.48	8.28	e
772305         AAAZ3680         8.71         4.709         4.85         1.82         1.82         1.83         0.88         7.22           8 247357         AAA23680         8.71         4.70         4.70         4.72         1.84         1.83         0.88         7.22           8 247357         AAA3548         0.64         0.04         0.05         4.45         1.86         4.47         1.83         1.88         3.84         1.83         1.88         3.84         1.83         1.88         3.84         1.83         1.88         3.84         1.83         1.88         3.84         1.83         1.88         3.84         1.83         2.45         4.65         1.47         1.49         6.57         1.83         1.88         3.84         1.83         4.45         1.89         8.88         8.74         4.85         1.88         1.89         8.88         8.74         4.85         1.86         8.88         8.88         1.81         4.48         1.89         8.88         8.74         4.85         1.87         4.48         8.89         8.74         8.88         1.83         1.81         4.89         8.88         1.89         1.89         8.88         1.83         1.89         1.89<	475	277598	N57002	1,48	£,	4.97	7.68	12.12	2.17	0.90	1.13	5.84	က
2.47587         AAAD35745         O.44         2.09         4.78         1.62         1.24         13.33         0.98         7.22           2.47587         NUSS138         0.01         0.05         4.64         1.96         14.29         1.91         4.85         3.94           2.47587         NUSS138         0.01         0.05         4.64         1.96         1.69         1.63         3.94         1.63         3.94         1.68         3.94         4.77         1.83         1.83         1.68         3.94         4.75         1.83         1.68         3.94         4.75         4.75         4.77<	<del>1</del> 78	773208	AA425850	9.71	47.09	4.85	1.82	1.90	6.84	12.13	2.88	3.52	က
2.475.57         NUSBTIAB         0.01         0.05         4.64         1.96         14.29         1.91         4.85         3.94           2.486.8         R1967.8         0.14         0.87         4.62         6.59         8.05         6.74         1.83         1.68           2.486.8         R1967.8         0.14         0.87         4.59         0.67         1.63         4.77         1.89         1.67           1.7720.4         AA423682         1.274         5.86         4.58         1.47         1.49         1.47         1.49         1.50         2.45           1.7700.4         AA423682         1.274         5.86         4.58         1.47         1.49         1.50         2.45         1.55           1.7700.4         AA42082         1.274         5.86         4.59         1.59         8.18         1.50         0.24         1.55         0.54         0.24         1.55         0.54	Ë	364865	AA035745	7.0	2.09	4.78	1.52	1.24	13,33	96.0	7.22	<b>4</b> .38	က
9.4886         FH 1876         CH 1         0.67         4.62         6.69         9.05         6.74         1.83         1.68           7.73264         A.425862         1.74         6.29         6.59         1.67         1.63         4.77         1.53         4.55         1.68           7.73264         A.4425862         1.74         58.08         4.56         1.47         1.49         5.57         13.00         2.45           7.73264         A.4425862         1.73         58.08         4.56         1.67         6.21         2.45         1.05         2.45           7.73264         A.4425862         1.73         3.33         4.62         7.28         8.84         0.39         0.24	178	247587	N58138	0.0	0.05	79.7	1.96	14.29	1.91	4.85	3.94	0.69	က
773264         7744264         7744264 <th< th=""><th>2</th><th>34888</th><th>R19878</th><th>2.7</th><th>0.67</th><th>4.62</th><th>6.69</th><th>9.05</th><th>6.74</th><th>1.83</th><th>1.68</th><th>2.65</th><th>e0</th></th<>	2	34888	R19878	2.7	0.67	4.62	6.69	9.05	6.74	1.83	1.68	2.65	e0
1         773204         AAA25862         1274         56.08         4.56         1.47         1.49         5.37         13.00         2.45           2         277227         N37483         0.02         0.09         4.50         1.67         6.21         2.55         8.18         1.05           3         770060         AA430545         7.38         33.38         4.52         7.28         8.45         6.19         0.54         0.26           1         51020         0.03         0.12         4.48         1.80         8.35         7.29         6.53         2.43           1         51020         0.03         0.12         4.49         1.80         8.36         6.53         0.24         0.29           2         28343         N2656         0.70         3.10         4.40         0.89         1.28         0.84         0.89         0.84         0.89         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89         0.84         0.89	5	72395	T51539	0.08	0.36	4.59	0.67	16.34	10.7	1.37	4.55	0.13	က
2 277277         NU37483         0.02         0.09         4.59         1.67         6.21         2.56         8.18         1.05           3 770068         AAA30546         7.38         33.38         4.62         7.28         8.45         6.19         0.54         0.28           4 6720         H19446         0.03         0.12         4.48         1.80         8.33         7.29         6.63         2.43           2 28343         N2556         0.70         3.10         4.44         8.62         9.86         4.53         0.64         0.64           2 28347         N2626         0.70         3.10         4.40         1.80         1.28         0.64         0.64         0.64           2 286347         N40256         0.70         3.10         4.40         1.80         1.28         0.64         0.	181	773204	AA425882	12.74	58.08	. 4.58	1.47	1.49	5.37	13.00	2.45	3,58	က
770066         AAA30545         7.38         35.38         4.62         7.28         8.43         6.19         0.54         0.26           1 51620         H19446         0.03         0.12         4.49         1.90         8.33         7.29         6.53         2.43           2 26343         N26556         0.70         3.10         4.44         862         9.86         4.53         0.84         0.84           2 268170         W02428         0.43         1.87         4.40         4.09         12.89         6.34         0.81         0.94           2 268170         W02428         0.43         1.87         0.28         4.34         0.89         1.80         0.81         0.94         0.94           2 268170         W02428         0.43         1.89         1.89         0.83         0.84         0.81         1.79         0.84         0.94         0.94           2 26827         AA122407         0.02         0.07         4.33         1.03         0.16         0.28         0.18         0.84         0.84         0.84         0.84         0.84         0.84         0.84         0.84         0.84         0.84         0.84         0.84         0.84         <	182	277327	N57483	90	0.09	95,4	1,67	6.21	2.65	8.18	1.05	7.68	က
6 F620         H19446         0.03         0.12         4.49         1.90         6.33         7.29         6.63         2.43           2 268343         N185556         0.70         3.10         4.44         8.62         8.86         4.53         0.64         0.94           2 268170         W02428         0.43         1.87         4.40         4.09         12.89         6.34         0.81         1.09           2 286170         W02428         0.43         1.87         4.40         4.09         12.89         6.34         0.81         1.09           2 286170         W02428         0.43         1.87         0.89         1.80         0.84         0.84         0.84           2 28626         A442570         0.06         0.22         4.33         1.035         0.16         0.26         2.13         0.69         0.84 <t< th=""><th>8</th><th>770068</th><th>AA430545</th><td>7.38</td><td>33.38</td><td>4.52</td><td>7.28</td><td>8.45</td><td>6.19</td><td>0.54</td><td>0.26</td><td>2.38</td><td>•</td></t<>	8	770068	AA430545	7.38	33.38	4.52	7.28	8.45	6.19	0.54	0.26	2.38	•
288343         NZBSSS         0.70         3.10         4.44         8.62         9.86         4.53         0.84         0.94           288170         WXZZZS         0.43         1.87         4.40         4.09         12.89         6.34         0.81         1.09           288170         WXZZS         0.43         1.87         4.40         4.09         12.89         6.34         0.81         1.09           1 2280547         AA128407         0.02         0.27         4.33         3.30         1.87         0.98         9.74         0.64           1 282056         AA424570         0.06         0.22         4.31         0.46         1.86         0.28         2.13         8.88           1 282057         AA484570         0.05         0.22         4.31         0.46         1.86         0.89         3.72         1.52           2 202058         MSP489         0.05         0.22         4.31         0.46         1.86         0.89         3.72         1.52         1.97         0.64         1.99         1.97         0.98         9.74         0.64         1.99         1.99         0.78         0.79         0.78         0.79         0.79         0.78	\$	51620	H19448	0.03	0.12	4.48	1.80	8.33	729	6.63	2.43	0.38	•
268770         W022426         0.43         1.67         4.09         12.89         6.34         0.81         1.09           728564         N86138         0.05         0.22         4.34         0.86         7.96         1.80         5.23         1.24           1 228567         AA128477         0.02         0.07         4.33         3.30         1.67         0.98         8.74         0.64           1 228268         AA484570         0.05         0.28         4.33         10.35         0.16         0.28         2.13         8.98           1 228268         N68498         0.05         0.22         4.31         0.46         1.86         0.28         2.13         8.98           2 205268         N68498         0.05         0.22         4.31         0.46         1.86         0.28         2.13         8.98           2 205268         N7566         0.61         2.26         4.28         7.32         8.76         0.78         0.78         1.01           2 205278         H7502         0.61         2.69         4.71         2.29         7.49         0.89         3.72         1.82         1.32         1.32         1.32         1.32         1.32	<b>3</b>	266343	N26556	0.70	3.10	7.	8.62	9.86	4.53	0.84	0.04	1.83	e
226567         N86139         0.05         0.22         4.34         0.86         7.96         1.80         5.23         1.24           1 52667         AA126407         0.02         0.07         4.33         3.30         1.67         0.98         9.74         0.84           9 09526         AA454570         0.06         0.28         4.33         10.35         0.16         0.28         2.13         8.98           9 09526         AA454570         0.06         0.22         4.31         0.46         1.86         0.88         3.72         0.54           1 250507         NF3643         0.42         1.81         4.28         7.32         0.76         2.49         3.67         1.81         1.01           1 250507         NF3643         0.61         2.18         7.29         2.40         5.57         1.28         1.52           1 67307         AA463373         4.86         2.04         4.10         5.29         2.40         5.57         1.28         1.37           2 84434         AA410187         2.86         1.16         2.04         4.10         2.89         2.40         5.89         1.37           2 84434         AA467864         0.04 <th>88</th> <th>296170</th> <th>W02428</th> <td>0.43</td> <td>1.87</td> <td>4.40</td> <td>4.09</td> <td>12.89</td> <td>6.34</td> <td>0.91</td> <td>1.09</td> <td>127</td> <td>m</td>	88	296170	W02428	0.43	1.87	4.40	4.09	12.89	6.34	0.91	1.09	127	m
526667         AA128407         CO.2         CO.7         4.33         3.30         1.67         CO.8         8.74         CO.64           9 09526         AA454570         G.06         0.28         4.33         10.35         0.16         0.28         2.13         8.08           2 282965         NUS489         0.05         0.22         4.31         0.46         1.86         0.26         2.13         8.08           2 205927         NY3663         0.42         1.81         4.28         7.32         0.76         3.67         1.01         1.01           2 205628         H57565         0.51         2.18         4.28         7.59         2.40         2.18         1.01           2 205628         H57565         0.01         0.06         4.11         5.2P         7.89         0.58         0.50         1.02           2 1022         4.12         4.12         3.45         0.81         2.40         0.58         1.37         1.02         1.37         1.02         1.03         1.34         1.37         1.02         1.34         1.34         1.37         1.44         1.05         1.06         0.68         0.58         0.50         1.06         1.34 <t< th=""><th>187</th><th>278504</th><th>N88139</th><td>0.05</td><td>0.22</td><td>4.34</td><td>0.98</td><td>7.96</td><td>1.80</td><td>5.23</td><td>1.24</td><td>8,83 53</td><td>က</td></t<>	187	278504	N88139	0.05	0.22	4.34	0.98	7.96	1.80	5.23	1.24	8,83 53	က
809526         AA454570         0.06         0.22         4.33         10.35         0.16         0.26         2.13         8.98           225266         N69489         0.05         0.22         4.31         0.46         1.96         0.28         3.72         15.21           225266         N7363         0.42         1.31         0.46         7.32         0.76         2.18         1.01           205826         N7363         0.42         1.81         4.28         7.32         9.76         2.18         1.01           205826         H57583         0.51         2.18         7.69         2.40         5.77         1.28         1.02           61302         2.44         2.28         2.40         2.28         0.58         0.58         0.58         1.37           28424         N64786         1.56         6.38         4.03         2.08         6.18         6.46         2.24         5.69           28424         N487385         1.06         4.28         4.03         2.50         4.10         2.28         7.47         8.01         2.44           28424         AA45786         1.06         4.28         4.03         0.29         4.10	<b>3</b> 5	526567	AA128407	0.02	20.0	4.33	3.30	1.67	0.98	9.74	9.64	9.67	eo
282968         N68469         0.05         0.22         4.31         0.46         1.86         0.88         3.72         15.21           295967         N73653         0.42         1.81         4.28         7.32         6.76         3.67         2.18         1.01           295828         H57585         0.61         2.16         4.28         7.58         2.40         5.77         1.28         1.01           611028         H57585         0.01         0.06         4.11         3.45         0.81         2.89         0.89         0.88         1.02           611028         AA485373         4.96         2.042         4.12         3.45         0.81         2.89         7.89         1.37           284624         N84786         1.58         6.38         4.03         2.08         6.18         2.44         5.89         1.37           2848247         AA487385         1.06         4.28         4.03         2.09         6.18         6.18         6.18         6.18         6.18         6.16         6.18         6.18         6.18         6.18         6.18         6.18         6.18         6.18         6.18         6.18         6.18         6.18         6.18	<b>8</b>	809526	AA454570	900	0.28	4.33	10.35	0.16	0.28	2.13	8.98	4.11	က
295807         N73653         0.42         1.81         4.28         7.32         9.76         3.67         2.18         1.01           205828         H57585         0.51         2.18         4.28         7.59         2.40         5.57         1.28         1.02           617028         H57585         0.51         2.18         4.28         7.59         0.53         0.90         0.68           611028         AA465373         4.96         20.42         4.12         3.45         0.81         2.08         7.59         0.58         0.90         0.68           611028         AA465373         4.96         20.42         4.12         3.45         0.81         7.47         8.01         2.44           284824         AA46786         1.53         6.38         4.03         2.09         6.18         6.48         2.24         5.69           841507         AA48738         1.08         4.28         4.02         10.98         2.50         4.10         3.36         1.15           280155         N47012         0.04         0.14         3.94         0.29         7.11         2.63         4.99           146878         AA463300         0.03 <t< th=""><th>\$</th><th>282968</th><th>N89489</th><th>0.05</th><th>0.22</th><th>4.31</th><th>0.46</th><th>1.86</th><th>0.68</th><th>3.72</th><th>15.21</th><th>3.80</th><th>60</th></t<>	\$	282968	N89489	0.05	0.22	4.31	0.46	1.86	0.68	3.72	15.21	3.80	60
205628         H57585         0.61         2.18         4.28         7.69         2.40         5.57         1.28         1.92           61397         749325         0.01         0.06         4.11         5.2P         7.69         0.68         0.90         0.68           811028         AA485373         4.96         20.42         4.12         3.45         0.81         2.68         7.75         1.37           289637         AA48738         1.68         11.62         4.06         3.07         1.98         7.47         8.01         2.44           284854         NA48738         1.68         4.28         4.03         2.09         6.18         8.46         2.24         5.69           280155         NA4078604         0.04         0.14         3.04         0.28         7.63         0.28         7.11         2.63         4.89           146678         AA487300         0.05         0.14         3.04         0.26         7.11         2.63         7.30         1.89         1.89         1.89           146678         AA443300         0.03         0.14         3.99         2.62         2.29         7.11         2.63         4.89           1	Ē	295997	N73563	0.42	1.81	4.28	7.32	9.76	3.67	2.18	1.01	1.74	60
67307         T49325         0.01         0.06         4.1n         5.2p         7.89         0.68         0.90         0.68           611028         AA485373         4.86         20.42         4.12         3.45         0.81         2.68         7.56         1.37           489637         AA485373         4.86         20.42         4.02         3.07         1.98         7.47         8.01         2.44           284824         N847786         1.56         6.38         4.03         2.09         6.18         6.48         2.24         5.69           841507         AA487385         1.08         4.28         4.02         10.98         2.50         4.10         3.36         1.15           280155         N47012         0.04         0.14         3.94         0.28         7.71         2.63         4.89           1466788         AA48300         0.05         0.75         2.29         7.71         2.63         4.89           146578         AA443300         0.03         0.10         3.89         3.64         1.89         1.89         1.83	192	205828	H57585	150	2.18	428	7.69	2.40	5.57	1.28	1.92	6.79	60
611028         AA465373         4.96         20.42         4.12         3.45         0.81         2.68         7.58         1.37           489637         AA4101878         2.86         11.62         4.06         3.07         1.98         7.47         8.01         2.44           284824         NA48738         1.56         6.38         4.03         2.09         6.18         8.46         2.24         5.69           81507         AA48738         1.06         4.26         4.02         10.88         2.50         4.10         3.36         1.15           280155         NA7012         0.06         0.14         3.94         0.26         7.71         2.63         4.89           1466788         AA483706         0.06         0.15         3.90         2.82         2.29         7.71         2.63         4.89           78459         AA44300         0.03         0.10         3.89         3.64         1.86         1.89         1.83	193	67397	T4832\$	0.01	90.0	4,10	5.28	7.89	0.58	06.0	0.68	8.45	•
4896.37         AA.101878         2.86         11.62         4.06         3.07         1.98         7.47         8.01         2.44           284824         NB4786         1.53         6.38         4.03         2.09         6.18         6.46         2.24         5.69           848247         AA487385         1.06         4.25         10.98         2.50         4.10         3.36         1.15           822497         AA487360         0.04         0.14         3.94         0.28         7.63         0.28         9.35         0.79           1466788         AA483730         0.46         1.79         3.90         2.62         2.29         7.11         2.63         4.89           1465788         AA443300         0.03         0.10         3.89         3.64         1.55         3.64         1.89         1.83	ğ	811028	AA485373	€.96	20.42	4.12	3.45	0.81	2.68	7.58	1.37	6.82	က
284624         N84786         1.53         6.38         4.03         2.09         6.18         6.46         2.24         5.69           841507         AA487385         1.06         4.26         4.02         10.38         2.50         4.10         3.36         1.15           822497         AA487385         1.06         0.14         3.94         0.28         7.63         0.28         9.35         0.79           280155         N47012         0.06         0.25         3.92         2.62         2.29         7.11         2.63         4.89           1468778         AA443300         0.03         0.10         3.89         3.05         1.55         3.84         1.89         1.83	195	489637	AA101878	2.86	11.62	4.06	3.07	1.98	7.47	8.01	2.44	1.39	က
841507         AA487385         1.08         4.28         4.02         10.98         2.50         4.10         3.36         1.15           882487         AA676604         0.04         0.14         3.94         0.20         7.63         0.28         9.35         0.79           280155         N47012         0.06         0.25         3.92         2.62         2.29         7.11         2.63         4.89           1466788         AAA83735         0.46         1.79         3.80         2.93         3.44         5.28         7.90         1.88           784589         AAA43300         0.03         0.10         3.69         3.05         1.55         3.84         1.89         1.83	<del>2</del>	284824	N84786	1.58	6.38	4.03	2.09	6.18	6.48	2.24	5.69	1.52	က
682487         AA676604         0.04         0.14         3.94         0.26         7.63         0.28         9.35         0.79           280155         NA7012         0.06         0.25         3.62         2.62         2.29         7.11         2.63         4.89           1466788         AAA83735         0.46         1.79         3.90         2.93         3.44         5.28         7.90         1.68           784589         AAA43500         0.03         0.10         3.69         3.05         1.55         3.64         1.89         1.83	197	841507	AA487385	1,08	4.28	4.02	10.98	2.50	4.10	3.36	1.15	2 2 3 3	60
280155         NAT012         0.08         0.25         2.62         2.29         7.11         2.63         4.89           1466768         AAA83736         0.46         1.79         3.90         2.93         3.44         5.28         7.90         1.68           784589         AAA43300         0.03         0.10         3.89         3.05         1.55         3.64         1.80         1.83	198	682497	AA676604	<b>5</b> 00	0.14	3.64	0.20	7.63	0.28	9.35	0.79	. <b>1</b> 63	n
1466788 AA483735 0.46 1,79 3.90 2.83 3.44 5,28 7,90 1.68 784589 AA443300 0.03 0.10 3.89 3.05 1.55 3.64 1,89 1.93	199	280155	N47012	90.0	0.25	3.82	25.52	2.29	7.11	2.63	4.89	4.00	<b>6</b>
784589 AA443300 0.03 0.10 3.89 3.05 1.55 3.64 1.88 1.83 1	82	1466788	AA883738	0.46	1.79	3.90	2.83	3.44	5.28	7.90	1.68	277	60
	50	784589	AA443300	0.03	0.10	3.89	3.05	1.55	3.64	1.86	1.83	11.30	n

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor semi-----

		Gen Rent		•	A	- AND THE PROBLEM OF	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F	old(MPM126/A	Semo
	IMAGE		AveBPH	AveTumor	Foid(AVe i umonz eBPH)	veBPH)	VeBPH)	veBPH)	VeBPH)	(Magru)		,
		Number				;	4 40	148	8.70	0.74	0.74	173
202	723972	AA292583	0.0	0.05	3.89	3.91	0/:/	900	0.35	1.33	8.10	က
5	858174	AA630620	9.0	0.11	3.89	6.41	6.16	2 6	0.63	0.88	3.47	n
3 2	782575	AA447522	0.38	1.39	3.88	8.55	10.7	<u> </u>	10.12	1.60	1.78	n
200	428168	AA001870	2.28	8.77	3.85	1.91	3.10	1 6	8.17	4.00	3.75	n
802	588326	AA130857	1.17	4.48	3.83	2.78	2.15 6.6	2 9	7.80	2.29	4.30	m
202	730838	AA417012	0.02	0.08	3.81	7.50	9 <del>4</del> .0	, t	2.48	6.88	2.62	က
208	837953	AA458578	7.85	29.78	3.79	3.76	28.0	7.8	68.89	5.74	6.57	m
209	503824	AA133959	0.12	4.0	3.78	1.62	90.0	2.5	1.69	0.76	1.61	60
340	208210	H85288	0.89	3.34	3.76	5.28	3.32	00 6	3.88	0.51	11.00	e
71	427996	AAD01841	0.1	0.43	3.75	3.08	90.0	75.0	4.24	12.61	4.50	en
212	123755	R01304	£.	6.90	3.75	0.50	97.0	3,5	8.23	5.08	2.80	e
213	438153	AA703249	14,14	52.21	3.69	1.02		2.57	2.67	3.32	4.58	n
214	143062	R71335	0.4	1.61	3.67	2.83	<b>t</b> 6	2.70	1.07	4.57	8.14	6
215	131589	R23727	0.16	0.60	3.65	5.05	2.39	1.72	4.91	6.33	5.22	e
218	240161	H82706	2.33	8.49	3.64	2.83	, i.	376	2.29	3.62	2.52	60
247	452116	AA707219	1,0	1.61	3.63	2.41	9 · · ·	, c	6.14	2.91	2,55	60
2,8	201125	R98472	1.05	3.81	3.61	3.83	4.00	2 2	4.29	1.20	2.78	e
218	128713	R07115	0.27	0.98	3,55	6.64	8 8	1.69	2.78	71.17	2.61	<b>6</b>
220	768179	AA424849	2.58	80.8	3.54	3.04	3.99	<u>6.</u>	3.77	2.33	2.51	က
2	250678	H96978	4.09	14.42	3.52	2.85		, Y	6.21	4.11	1.47	m
122	273489	N33263	60.0	0.32	3.52	3.88	2.6/	 	7.82	1.94	1.82	m
223	766785	AA496455	1.95	8.84	3.51	2.12	3.80	Ç •	1.7	3.01	7.18	က
ž	815047	AA465166	2.68	9.34	3.49	2.62	7.60	7 1.	7.25	2.34	2.03	n
22	384670	AA709333	4.17	14.53	3.48	3.11	- 6.4	, c	3.5	2.28	3.28	6
ង	782200	AA431964	1.53	5.30	3.47	2.48	9.95	104	6.25	3.48	2.49	n
ä	62730	H28500	9. 2.	22.84	3.45	2.33	- 78 c	99.6	6.68	4.33	2.05	n
228	125311	R05810	1.92	6.62	, 1	1.07	, t. c.	7.87	1,10	0.49	1.77	ė
877	770830	AA434404	8.05	27.68	3.44	6.26		2.58	5.27	5.89	1.71	c
23	796268	AA480827	0.55	1.89	3.43	1.43	5.00 7.00	7.21	4.18	1.12	1.39	<b>6</b>
231	204148	02(_)	2.07	7.07	3.42	2.81	2.0		3.86	08.0	08.0	n
23	611373	AA176833	0.01	9.	3.41	9.87	5	5.78	5.41	1.13	5.09	t)
23	325380	AA284285	0.01	0.03	3.40	1.80	50,	7	1.12	1.58	6.01	n
ន្ត	434822	AA703114	0.80	2.72	3,38	0°.0	4.75	2.95	4.55	1.19	ą į	es
22	1048033	AA778876	1.87	<b>8</b> .35	3.38	8.4	,					

	IMAGE Clone ID	Gen Benk Accession Number	AveBPH	AveTumor	Fold(AvoTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor eBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH)	Fold(MPN120/A veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A   vaBPH)	Fold(MPM122/A F veBPH)	-old(MPM123/A I	Fold(MPM125/A veBPH)	# of Tumor Samples
238	882483	AA676598	2.76	9.30	3.38	1.34	4.86	2.99	5.61	3.78	1.72	
237	767130	AA424535	0.49	1.67	3.38	1.02	0.74	1.14	3,10	10.17	4.12	ဗ
238	128321	R06479	0.03	0.10	3.38	2.12	4.4	3.73	7,41	0.36	221	6
238	230440	H75328	0.07	0.22	3.37	1.97	3.57	3.44	2.89	8.21	0.16	ဇာ
240	824271	AA181179	6.78	22.81	3.36	3.53	3.85	2.87	5.02	2.93	1.98	၈
7	123265	R00284	2.30	7.72	3.35	4.22	9.13	4.42	99'0	0.86	0.68	ღ
242	357236	W83638	0.72	2.40	3.35	5.76	2.69	1.43	1.48	3.63	5.19	ო
243	123331	R00479	1.03	3.43	3.33	2.37	4.72	5.66	0.98	3.48	2.80	<b>6</b> )
<b>7</b>	240691	H78134	1.85	6.17	3.33	2.79	3.07	1.59	7.73	3.58	121	<b>m</b>
245	347182	W80811	0.02	0.07	3.32	0.46	5.03	5.53	<b>16.4</b>	2.81	1.19	8
240	25085	H23232	0.03	0.11	3.31	1.94	4.36	6.28	4.13	2.87	0.31	ო
247	160828	H25019	0.41	1.38	3.30	2.02	4.53	3.17	4,45	2.72	2.83	ო
248	1505294	AA 205838	1.3	3,69	56.6	c. 0.	2.47	7.2*	1.55	3.28	3.34	ღ
348	282053	H98912	7.69	25.26	3.28	2.38	3.42	3.48	6.75	1.98	1.68	E)
250	741067	AA478436	17.59	. 57.78	3.28	2.99	3.39	3.14	2.34	2.25	5.58	က
251	490178	AA121288	3.50	11.48	3.28	2.64	4.34	2.28	4.85	1.83	3.72	6
252	208741	H83175	0.09	0.30	3.28	2.00	4.61	3.24	2.64	2.82	4.27	6
283	418811	W86468	3.63	11.85	3.20	2.01	2.14	2.13	5.36	4.86	3.07	က
52	416092	W85881	0.50	1.62	3.28	2.00	2.31	1.93	3.05	3.65	6.62	<b>6</b>
255	221581	H82525	0.55	1.79	3,26	1.18	1.35	3.60	1.43	6.94	5.05	<b>6</b> 0
<b>5</b> 2	223274	H86518	0.05	0.17	3.25	3.24	0.19	0.18	2.95	5.14	7.78	6
257	44387	H06525	1.17	3.78	3.24	6.67	2.69	4.15	1.74	3.08	1.09	67
<b>728</b>	110787	T90839	0.25	0.80	3,23	2.49	3.52	2.19	3.80	2.05	5.34	60
220	121948	197762	4.17	13.48	3.23	1.59	4.12	2.42	5,41	3.80	2.03	6
<b>5</b> 80	782812	AA448251	0.08	0.18	3.22	4.87	5.08	1.22	2.83	3.17	2.10	n
<b>58</b>	145513	R77718	0.67	2.18	3.22	1.90	4.01	2.67	3.28	2.99	4.46	<b>m</b>
	435318	AA699931	6.14	19.70	3.21	2.30	5.27	1.85	3.83	2.74	3.25	9
	210698	H66877	1.41	4.62	3.21	1.58	2.24	3.47	5.80	5.20	0.65	6
	197067	R83401	0.38	1.18	3.19	1.88	1.56	3.78	3.20	2.11	6.62	<b>6</b>
	418374	W86202	3.18	10.13	3.19	4.89	3.63	2.43	2.00	3.28	2.91	e
_	1422794	AA827405	2.02	6.43	3.18	2.60	3.91	2.60	5.88	1.06	3.06	е
	137017	R35865	0.03	0.08	3.18	1.47	4.27	1.59	8.08	3.23	0.45	en
		N34117	0.21	9.68	3.18	2.32	3.85	2.53	3.68	3.77	2.82	•
7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7	433253 /	AA699427	6.81	17.79	3.17	4.18	4.57	2.43	1,60	1.72	4.54	•

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	100	Gen Bank			Fold(AveTamor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumo	Fold(MPM1201A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A veBPH)	Fold(MPM123/A	Fold(MPM12&A veBPH)	s of Tumo Samples
		Accession	AveBPH	Averumor	•BPH)	veBPH)	Vebruj		•	ç	6.79	က
		Number				376	3.26	2.34	2.32	2,7	· ·	
270	1475734	AA872692	7.39	23.26	ن. د :	5 4	6.62	3,35	1.38	3.88	81.2	
7	268876	N28028	0.51	1.59	3.14	<b>Q</b>		90 8	1.33	2.11	2.80	<b></b>
	70m2	TR1899	15.17	47.49	3.13	3.66	76.4	) ¥	2.35	3.41	4.13	eo
; [	71623	A4477978.	1.12	3.51	3.13	2.45	3.57	6.43	76 40	4.36	3.37	က
	844404	A A A R S. S. T. B.	2.17	8.72	3.09	1.06	2.45	0,1	121	5.53	2.34	m
<b>* !</b>	201710	A04614	0.66	2.04	3.09	3.84	3.31	2.13	<u> </u>	1.38	1.27	က
9	17077	AA702480	2.85	8.16	3.08	3.60	2.77	\$ 3	פ	3.43	2.39	m
<b>\$</b>	7 1000	AAABARRY	870	1,48	3.07	2.06	2.12	3.31		1.77	4.45	က
<b>5</b> i	1000	A & B B D D D D	6 6	1.47	3.05	1.65	4.32	232		4	3.82	65
817	431280	PA002031	1 92	5.87	3.05	2.21	3.82	1.69	 	2.28	3.78	က
278	*70*LZ	7471/14		1.50	3.04	4.39	4.19	4.69	G 5	5.48	1.81	က
280	743323	AAAOOO	, ,	88.85	3.6	1.01	1.73	3.23	à.	; <del>,</del>	3.69	m
281	80050	163245	7 17	2 2	5	2.12	4.97	1.27	4.24	7.	808	က
282	. 1500162	AA886739	80.7	2 6	3.03	2.45	3.22	6.41	2.32		7	87
283	268234	N28882	1.72			2.89	4.10	3.83	3.02	2.70		· •
<b>78</b>	100623	R96561	0.51	<u> </u>	3.5	3 7	3.80	2.86	4.40	3.55	8 :	, (
285	41305	R56877	19.83	60.04	3.03	- 4° 6	¥	2.63	16.4	4,1	1.32	<b>o</b> (
288	795555	AA459679	1.63	7.87	3.03	8 1	00.4	2.37	4.02	3.51	2.42	m (
287	416808	W86779	0.57	1.71	3.02	1.11		325	4.15	4.88	1.97	ro
288	242698	H94262	1.10	3.50	3.00	1.70	7.77	4 22	3.89	4.83	2.18	60
280	384352	AA022496	0.03	0.08	3.00	2.52	0.39	1 88 1 7	2.22	3.28	5.43	က
8	244050	N34042	0.08	0.17	3.00	1.60	A	) - «	3.80	2.68	2.58	0
ğ	544639	AA074877	1.05	3.15	2.00	1.72	3.6	98.6	88.	2.63	0.92	60
202	862620	AA683058	17.95	\$3.52	2.98	3.39	3.07	. ¥	5.82	1.07	17.4	60
8	797000	AA463500	0.11	0.33	2.98	<u>.</u> 2	3.21	3 2	3.28	1.52	2.20	೯
ž	1483252		18.18	\$4.08	2.98	4.82	5.55	2 42	20.	3.10	\$	က
295	447418		1.15	3.41	2.98	5.58	i c	4.53	1.90	1,93	3.19	e0
298	1475308	_	7	4.58	2.98	2.82	3.54	98.6	2.74	3.89	2.47	<b>.</b>
297	1913		0.22	0.65	2.95	.35 .35	, c. c.	171	2.34	0.80	3.84	•
88	1456937	_	1.15	3.38	2.87	5.57	9.30	2.0	3.06	2.04	3.43	•
288	201039		0.83	2.45	2.93	2.78	6.80 1.80		1,10	1.08	0.29	63
90	283769		0.75	2.19	2.93	3.82	6.6	, e	0.21	77.7	2.22	ю
Š	427930	•	0.05	0.14	2.91	0.89	4	3 4 6	2.07	3.85	1.25	69
302	307244		0.35	<u>5</u>	2.91	270	4.84	: ;	10.0	4.41	<u>₹</u>	n
g	810843			0.17	2.90	 ¥.	6.13	<u> </u>				
							c					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	MAGE Clone ID	Gen Bank Accession Number	AveBPH	AveTumor	Fold(Avetumoriay Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A \$ of T •BFH) veBFH) veBFH) veBPH) Sam	Fold(MPM120/A   vebPH)	Fold(MPM1211A veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A   VBBPH)	Fold(MPM123/A ) veBPH)	Fold(MPM125// veBPH)	Sam.
Š	1480820	•	3.62	10.49	2.90	2.35	2.11	3.28	1.57	3.43	4.63	m
8			0.87	2.50	2.89	3.08	3.41	3.69	2.10	2.31	2.7	es
8	366154	AA062813	0.01	0.03	2.85	3.87	3.77	1.00	1.23	1.00	6.23	C
307	282104	N51498	0.93	2.66	2.84	2.66	4.75	3.79	1.52	0.39	3.87	က
88	344308	W70147	0.46	1.31	2.81	3.18	3.47	5.50	1.78	1.21	1,74	e,
808	;	H08725	2.19	6.17	2.81	2.20	3.24	3.18	3.52	2.33	240	***
310		R92435	90.0	0.11	2.80	2.56	4.53	3.04	2.48	3.30	0.85	•
3		N25262	1.59	4.40	2.77	3.68	3.24	2.59	3.41	1.52	2.19	₹,
312	757108	AA498133	0.65	1.78	2.78	3.73	4.67	3.45	124	1.90	1.53	<b>r</b> >
313	204122	H61886	1.31	3.59	2.75	2.36	4.23	3.40	3.07	2.93	0.52	•,
314	838545	AA489791	0.42	1.1	2.73	5.37	0.89	1.38	1.18	3.64	3.82	۲,
315		AA454702	0.02	0.05	2.70	9.54	0.56	3.30	2.89	4.45	4.47	••
348		R08362	0.30	0.81	2.69	2.13	3.70	2.31	1.13	3.58	3.32	۲,
317		N51563	0.97	2.62	2.69	1.61	3.40	1.91	3.26	3.22	272	**
318	758284	AA04231	0.78	2.08	2.67	1.82	2.31	1.16	3.46	3.11	4.17	••
319	150314	HD0817	2.18	5.01	2.67	3.69	3.80	3.23	2.04	1.07	2.09	••
320	288938	N62696	9.0	0.11	2.67	1.59	3.47	3.00	0.69	2.99	3.87	۴,
321	757440	AA437228	0.05	0.13	2.67	0.21	4.19	1.79	4.52	0.21	5.08	E.)
ğ	178554	H45289	0.87	2.28	2.64	1.89	2.02	3.72	3.33	3.56	1.30	62
g	296057	N69989	99.0	1.79	2.62	2.03	3.01	2.54	1.85	3.26	3.04	63
33	755289	AA486359	1.69	4.39	2.60	4.47	0.52	1,93	3.99	0.70	3.88	67
325	725321	AA291749	0.87	2.27	2.59	3.21	3.07	3.61	2.30	1.62	1.73	<b>6</b> 0
33	1467631	AAB83431	1.17	3.02	2.50	1.16	1.22	4.32	0.72	3.58	4.52	60
321	120097	T95151	0,60	<b>2</b> .	2.58	2.09	2.01	1.14	3.86	3.01	3.31	47
328	159455	H15748	0.02	0.0	2.58	0.62	3.70	2.31	0.64	3.72	4.51	۳,
33	1460688	AA868802	6.57	16.82	2.58	1.39	3.08	<u>r</u> .	2.32	3.41	3.22	67
88	757388	AA437128	0.02	90:0	2.55	3.92	2.20	3.37	0.77	0.24	<b>4</b> .83	m
뚔	450612	AAB82452	27.0	1.89	2.55	1.17	0.96	0.92	3.00	4.79	7.7	е
ä	342543	W68141	0.02	5,0	2.53	4.17	4.4	0.58	1.17	4.50	0:30	n
g	503741	AA131468	1.05	2.65	2,53	3.31	3.85	3.58	1.37	4.19	1.85	67
ន្ត	784130	AA432075	17.73	44.77	2.52	3,17	4.60	0.70	1.58	3.28	1.78	69
8	325370	W52208	0.03	0.08	2.51	3,68	3.87	2.92	1,13	0.31	3.14	n
2	288683	N62394	0.78	1.95	2.50	3.07	3.98	0.85	1.90	1.37	3.87	ro
ğ	344359	WT3377	0.07	0.18	2.46	0.90	3.52	0.14	3.08	7.02	0.08	67
						#	10					
						,	•					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

Members         114         278         241         334         140         123         133           Members         114         278         241         316         319         152         165         319         314           Moscost         0.01         0.35         2.41         316         313         122         165         313           MOSCOST         0.02         2.31         0.06         4.21         100         319         314         314           MOSCOST         0.03         2.31         1.06         3.23         1.00         3.13         3.14         3.15         3.14         3.15         3.14         3.15         3.14         3.15         3.14         3.15         3.14         3.14         3.14         3.14         3.14         3.14         3.15         3.14 <th< th=""><th>_</th><th>IMAGE</th><th>Gen Bank</th><th>Magaria</th><th>AveTimor</th><th>Fold(AveTumon/Av Fold(MPM120)A Fold(mPM120)A VeBPH) veBPH) veBPH) veBPH) veBPH)</th><th>Fold (Mr. 14 14 14 14 14 14 14 14 14 14 14 14 14</th><th>VaBPH</th><th>veBPH)</th><th>VeBPH)</th><th></th><th></th><th>•</th></th<>	_	IMAGE	Gen Bank	Magaria	AveTimor	Fold(AveTumon/Av Fold(MPM120)A Fold(mPM120)A VeBPH) veBPH) veBPH) veBPH) veBPH)	Fold (Mr. 14 14 14 14 14 14 14 14 14 14 14 14 14	VaBPH	veBPH)	VeBPH)			•
SSSPRI (NECKAL I.16)         1.77         2.41         3.60         3.67         3.34         1.60         3.55         1.50         3.55         1.50         3.55         1.50         3.55         1.50         3.55         1.50         3.55         1.50         3.55         1.50         3.55         1.50         3.55         1.50         3.55         3.55         3.25<	- 0	Ol enol:	Accession	AVebra		•8PH)	Vebru			9	123	1.35	•
CORREST         AVAILAGE         2.7         3.1         3.1         1.2         1.65         2.7         3.1         3.2         3.2         1.6         3.1         3.2         3.2         1.6         3.1         3.2         <	'		Number	!		*** 6	3.60	3.67	33,4	2 ·		50.50	က
CASES         AVEOREM         0.14         0.35         2.40         3.17         0.25         0.05         0.25           252891         AVEOREM         0.02         2.31         0.06         4.21         1.00         1.22         3.01         1.39           252891         AVEOREM         0.08         0.23         2.30         0.24         1.00         1.25         3.01         1.39           452895         AVEOREM         0.08         0.20         2.31         0.06         0.21         1.20         3.09         0.19         3.19         3.11         1.39           650848         AVEOREM         0.08         2.15         1.20         2.42         1.00         1.25         3.01         1.39           120048         AVEOREM         0.03         1.42         0.03         1.42         0.04         2.04         3.01         1.39         0.04         0.04         2.04         3.01         1.00         1.30         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04         0.04 <td></td> <td>53391</td> <td>R16241</td> <td>1.16</td> <td>2.79</td> <td>- ·</td> <td>4</td> <td>3.19</td> <td>1.82</td> <td>1.65</td> <td>B) :</td> <td>;</td> <td>e</td>		53391	R16241	1.16	2.79	- ·	4	3.19	1.82	1.65	B) :	;	e
225821         NGORS         0.62         1.49         2.40         3.50         4.51         1.00         1.52         3.40         3.51         3.60         4.51         3.60         4.51         3.60         4.51         3.60         4.51         3.60         3.71         3.60         3.71         3.60         3.71         3.60         3.71         3.60         3.71         3.60         3.71         3.60         3.71         3.60         3.71         3.60         3.71         3.72         3.71         3.72         3.72         3.71         3.72         3.71         3.72		42B823	AA009803	0.1	0.35	2.40	2 6	123	0.26	4.12	0.85	26.3	•
255587         X77368         0.08         0.20         2.31         0.80         3.21         1.70         319         301         1.71         319         301         1.71         319         301         1.71         319         301         1.71         319         301         1.71         319         301         1.71         319         301         1.71         319         301         1.71         318         301         1.71         302         4.22         300         0.74         0.24         3.20         0.24         3.20         0.24         4.85         7.71         4.85         7.72         0.24         3.00         0.04         4.85         0.74		0KR242	N30855	0.62	1.49	2.40	3.6	) ;	9	1.52	3.40	3.14	3
685779         AATJONT 22         CAS         2.43         2.44         1.10         3.57         0.38         2.58         2.43         3.24         1.10         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.38         3.57         0.39         3.57         0.39         3.57         0.39         3.57         0.39         3.57         0.39         3.57         0.39         3.57         0.39			A17790	0.09	0.20	2.31	0.60	17.4	,	3.19	3.01	1.39	m
455084   AAA77725   Use   AAA777208   AAA777208   AAA777208   AAA777208   AAA777208   AAA777208   AAA777208   AAA777208   AAA777708   AAA777708   AAA777708   AAA777708   AAA777708   AAA777708   AAA777708   AAA77708   AAA7708   AAA77708   AAA77708   AAA7708   AAA77708   AAA7708   AAA7708   AAA7708   A		/80CC	N21300		200	2.30	1.29	3.24	2.		0.38	3.18	e
SESTIPA         AMAZIGNITO         20.69         48.64         2.55         1.00         3.87         0.24         3.00         0.24           178506         RODINIS         0.04         0.09         2.12         4.12         3.45         0.42         3.04         0.08           1280704         AMASINES         0.04         0.09         2.12         4.12         3.45         0.42         3.04         0.08           1280704         AMASINES         0.01         0.18         13.25         0.74         0.74         1.89         0.74         4.48           500815         AAVISORTO         0.02         1.10         0.02         1.10         0.02         2.14         0.74         1.89         0.74         4.48         1.74         4.48         0.04         2.74         1.89         0.74         4.48         1.74         4.48         0.04         2.14         4.48         0.04         2.14         1.74         1.89         0.74         1.74         4.24         1.74         4.24         1.74         4.24         1.74         4.24         1.74         4.24         1.74         4.24         1.74         4.24         1.74         4.24         1.74         4.24		452068	AA707125	9 9 9	3		2.43	3.80	0.18	6.5		4 04	e
18856   R10154   Clot   Clot		585779	AA135870	20.69	48.64	2.25	9 6	2.87	0.24	3.30	0.24	9.	• •
699048         AAA45700         0.26         2.12         4.72         2.74         2.38         1.76         4.85         7.7         2.38         1.70         4.85         7.7         2.38         1.70         4.85         7.7         2.38         1.70         4.85         7.7         2.38         1.70         4.85         7.7         2.84         8.7         2.44         8.3         2.44         8.3         1.7         4.85         7.7         2.8         2.7         1.89         0.74         4.85         7.7         4.85         7.7         4.80         0.7         4.85         7.7         4.80         0.7         4.80         0.7         4.85         7.7         4.80         0.7         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         4.80         0.7         0.7         0.7		128508	R10154	ŝ	0.09	2.15	02.L		0.42	3.04	0.88	0.82	<b>,</b>
200763         Avidential states         0.50         14.36         0.50         2.77         Case         0.74         4.86         0.74         4.86         0.74         4.86         0.74         1.87         1.87<			44407808	91.0	0.76	2.12	4.12	0 ·	5 6	178	4.85	73.83	~
267075 NGSSSS         AMSSSS         COLOR OF SERVICES         COLOR OF		200	Appendix		2 6	14.36	0.58	2.77	7.30		47 48	32.88	~
BODISS         AATISSIES         CAG         1.20         1.40         0.82         3.37         1.30         1.40           \$60101         AAGSCAZZ         0.02         0.22         11.30         1.40         0.82         2.36         49.00         1.07           \$61101         AAGSCAZZ         0.02         0.20         1.30         1.40         0.23         2.26         49.00         1.07           \$61701         AAGSCAZZ         0.02         0.20         0.20         1.00         0.20         0.20         1.00         0.27         0.26         1.11         2.02         1.11         2.02         1.11         2.02         1.11         2.02         1.11         2.00         1.1		280763	N20263	3	3 5	43.05	0.74	0.74	 8	£ 7.0	er c	58.40	7
951101         AAGZGATZ         0.02         0.13         1.6         2.93         2.36         49.00         1.07           977810         AAGZGATZ         0.02         0.13         0.02         1.60         0.16         2.36         2.93         2.87         34.78           977810         AAGZGATZ         0.04         0.05         1.00         1.00         1.60         0.16         2.36         2.87         3.71         1.11         2.77         1.46         0.20         0.20         0.24         1.11         2.74         4.85         0.24         3.71         1.11         2.74         4.85         0.24         3.71         1.11         2.74         4.85         0.24         3.71         1.11         2.74         4.85         0.24         3.71         1.11         2.74         4.85         0.24         3.71         1.17         2.74         4.85         0.24         3.71         1.17         2.74         4.85         0.24         3.71         1.17         2.74         4.85         0.24         3.71         1.17         2.74         4.85         0.24         3.71         1.17         2.74         4.85         0.24         3.71         1.17         2.74         4.85 <td></td> <td>590853</td> <td>AA158162</td> <td>9</td> <td>0.70</td> <td></td> <td>071</td> <td>0.82</td> <td>3.37</td> <td>1.38</td> <td>0 1</td> <td></td> <td>~</td>		590853	AA158162	9	0.70		071	0.82	3.37	1.38	0 1		~
377810         AAADSSSSS         0.13         1.34         10.02         1.40         0.16         2.38         2.67         34.78           67787         H16833         0.06         0.50         7.39         1.60         0.16         2.38         2.67         34.78           437884         R83055         0.29         2.25         7.79         8.43         33.82         0.37         0.58         1.11           58586         AABSSZZZ         0.12         0.49         7.74         1.97         2.74         4.85         0.24         35.70           58586         AABSSZZZ         0.12         0.49         7.44         1.97         2.74         4.85         0.24         35.70           43017         ALGIOZZ         0.10         0.75         1.46         0.70         1.19         3.74         1.47         35.70         0.48         1.47         35.70         0.48         35.70         0.48         3.74         4.85         0.24         35.70         0.48         3.74         4.85         0.74         1.47         37.80         0.48         1.47         33.12         2.43         1.47         35.70         0.48         1.47         33.12         2.43		951101	AA620472	0.02	0.22	J. J.	2 4	203	2.38	49.00	1.07	0.0	
60787         H16833         0.08         0.50         7.90         1.60         0.10         0.11         0.12         0.11		377810	AA055833	0.13	¥.	10.02	0.40	9,0	2.38	2.67	34.78	28.6	<b>y</b>
137884         R83085         0.29         2.25         1.79         84.3         33.82         0.29         3.24	*	2000	LYARTS	0.06	0.50	7.90	1.60	9 9	6	0.58	1.1	2,4	~
1384   1450   146   146   146   146   146   147   146   147   146   147   146   147   146   147   146   147   146   147   146   147   146   147   146   147   146   147   146   147   148   147   148   147   148   147   148   147   148   147   148   147   148   147   148   148   147   148   14		20100	200011	0,0	2.25	7.79	8.43	33.82		67.0	32.14	1,71	7
B55568         AA688222         0.14         1.97         2.74         4.55         0.24         1.77           358900         WA2283         0.06         0.48         7.64         1.97         2.74         4.55         0.24         1.77           358900         WA2283         0.06         0.48         7.64         1.97         2.73         2.935         1.18         3.78         1.47           45012         A.010225         1.19         8.16         6.81         1.533         1.931         2.50         1.64         0.68         0.46         0.68         0.46         0.68         0.41         0.55         1.01         0.68         0.41         0.68         0.57         1.64         0.68         0.41         0.68         0.41         0.68         0.41         0.68         0.41         0.68         0.41         0.68         0.41         0.68         1.01         0.68         0.43         1.07         0.12         0.12         1.01         0.68         0.43         1.01         0.68         0.43         1.01         0.68         0.43         1.01         0.68         0.43         1.01         0.68         0.41         0.68         0.41         0.68         0.43 <td>_</td> <td>13/984</td> <td>K83083</td> <td></td> <td>70</td> <td>1.77</td> <td>1.46</td> <td>0.70</td> <td>B.</td> <td>6</td> <td>25.70</td> <td>99.0</td> <td>~</td>	_	13/984	K83083		70	1.77	1.46	0.70	B.	6	25.70	99.0	~
358990         W42283         0.06         0.74         7.77         33.12         2.43         1.19         3.78         1.47           420182         AA010228         0.10         0.75         7.47         7.32         29.35         1.19         3.78         1.47           15131         150111         0.10         0.75         7.47         7.32         29.35         1.05         0.48         0.46           15131         150111         0.47         3.19         6.81         1.37         22.88         0.57         1.64         0.68           152028         142802         0.22         1.66         6.89         1.07         0.12         0.12         0.52         1.01           42808         AA026405         0.08         0.23         6.48         1.07         0.12         0.02         1.64         0.68           42808         AA026405         0.08         0.23         6.48         1.07         0.12         0.12         1.04         0.55         1.01           42808         AA026406         0.02         0.22         6.48         0.41         0.12         0.12         1.03         1.04         0.55         1.01         0.12         1.0	~	858585	AA689222	3	; ;	181	1.97	2.74	4.55	* * * * * * * * * * * * * * * * * * * *	,	284	~
450162         AAD10228         0.10         0.75         7.47         7.32         2.935         1.05         0.48         0.46           151477         H02837         1.19         8.16         6.81         7.32         2.935         1.05         0.68           151477         H02837         1.19         8.16         6.81         15.33         1.01         0.68           52026         H22032         0.25         1.68         6.89         1.37         2.286         0.57         1.64         0.68           62026         H22032         0.25         1.68         6.89         1.37         2.286         0.57         1.01         0.62         1.01           428086         0.02         0.32         6.48         0.49         0.41         0.39         1.36         1.39         1.01           428086         0.02         0.27         1.39         6.27         0.41         0.39         1.36         1.30	_	358990	W92283	90.0	0 4 0	<b>5</b> !	23 13	2.43	1.19	3.78	1.47		•
151477         H02837         1.19         8.16         6.87         1.34         1931         2.60         1.64         0.68           81331         780111         0.47         3.19         6.81         15.33         1931         2.60         1.64         0.68           82026         H22832         0.25         1.69         6.89         13.67         22.88         0.67         1.63         1.01           42008         H22832         0.69         0.53         6.43         1.07         0.12         0.32         1.84         1.84         1.84         1.01           42008         AA08142         0.22         1.39         6.27         0.88         0.41         0.36         1.35         1.84         1.35         1.01         0.32         1.84         6.88         0.41         0.32         1.84         6.88         0.41         0.36         0.35         1.84         0.85         1.30         0.85         1.30         0.85         1.30         0.85         1.30         0.85         1.30         0.85         1.30         0.85         1.30         0.81         1.30         0.85         1.30         0.85         1.30         0.81         1.30         0.85	J	430192	AA010328	0.10	0.75	. 13.1	4.35	20.35	1.05	0.48	0.48	25 T	• (
81331         T60111         0.47         3.19         6.81         13.33         1.01         1.01         1.01           62028         H22832         0.25         1.68         6.89         13.67         22.88         0.57         1.63         1.01           62028         H22832         0.25         1.68         6.89         1.07         0.12         0.52         31.38           429086         AAA086405         0.08         0.53         6.48         6.88         2.35         1.00         0.55         1.30         1.30           284449         NZ1233         0.06         0.32         6.48         6.88         2.35         1.00         0.55         1.30         1.30           284449         NZ1233         0.06         0.32         6.48         2.04         2.77         1.77         2.77         1.30 <td>47</td> <td>151477</td> <td>H02837</td> <td>1.19</td> <td>8.18</td> <td>6.87</td> <td>26.7</td> <td>40 34</td> <td>2.50</td> <td>1.64</td> <td>0.68</td> <td>1.40</td> <td>٠ (</td>	47	151477	H02837	1.19	8.18	6.87	26.7	40 34	2.50	1.64	0.68	1.40	٠ (
62028         12612         6.89         1357         2200         136         146         146	. «	81331	T80111	0.47	3,18	6.81	15.33		0.57	1.63	1.01	0.38	7
659627         AA606405         0.08         0.53         6.43         1.07         0.12         0.12         1.30         1.35         18.48           429087         AA606405         0.08         0.22         1.39         6.27         0.88         0.41         0.36         1.35         1.30         2           429088         AA005168         0.22         1.39         6.48         6.98         2.35         1.09         0.55         1.30         2           264449         N21233         0.06         0.32         6.48         6.98         2.37         1.77         2.77         1.30         2           764368         AA436142         8.25         44.98         5.45         2.04         2.77         1.77         2.77         1.30         2         1.30         2         1.30         2         1.30         0.91         0.92 <td< td=""><td>, ,</td><td>RCOCA BCOCA</td><td>H22832</td><td>0.25</td><td>1.68</td><td>6.89</td><td>13.67</td><td>7.00</td><td>1 1 2</td><td>0.52</td><td>31,38</td><td>5.38</td><td>7</td></td<>	, ,	RCOCA BCOCA	H22832	0.25	1.68	6.89	13.67	7.00	1 1 2	0.52	31,38	5.38	7
CARSAN         AAACASSIGN         0.22         1.39         6.27         0.88         0.41         0.30         1.30         2.25         1.30         1.30         2.24         1.30         0.55         1.30         2.30         1.30         2.24         1.30         0.55         1.30		04040	AABBAADS	0.08	0.53	6.43	1,07	21.0		1.35	18.48	16.17	~
428088         AMODSTURE         0.22         6.48         6.98         2.35         1.08         0.33           264149         NZ1Z33         0.06         0.32         6.48         6.98         2.77         1.77         2.77         1130           764368         AA438142         8.25         44.96         5.45         2.04         2.77         1.77         2.77         1130           76826         AA42800         0.06         0.31         5.18         1.16         22.31         1,49         3.68         0.91           787166         AA42800         0.06         0.31         5.19         0.47         1,51         0.96         1,45         0.13           303139         N90806         0.52         2.71         5.17         4.05         23.41         0.96         1,45         0.13           757166         AA43850         0.03         0.16         5.09         0.47         1,51         0.94         20.25         2.06           247103         N53859         0.05         0.22         4.82         3.70         1,71         1,84         0.22         2.40           234376         AA43811         0.04         0.21         4.73         <	9	70000	COLOGO		130	6.27	0.88	0.41	0.0	9 6	1.30	20.60	7
264449         RZ1233         0.00         0.22         2.04         2.77         1.77         2.11         2.11           76436         AA436142         8.25         44.86         5.45         2.04         2.77         1.77         2.11         0.91           76826         AA43800         0.06         0.31         5.18         1.16         22.31         1,49         3.88         0.91           303139         N90806         0.52         2.77         5.17         4.05         23.41         0.96         1,45         0.13           377105         AA43850         0.03         0.16         5.09         0.47         1,51         0.94         20.25         2.06           247103         N53959         0.03         0.16         5.09         0.47         1,71         1,84         0.22         2.88           247103         N53959         0.05         0.22         4.82         3.70         1,71         1,84         0.22         2.88           234376         AZ3288         2.94         14.18         4.82         3.75         1,45         2.24         3.15         2.40           247103         AZ3288         AA482119         0.04 <t< td=""><td></td><td><b>429088</b></td><td>AA005186</td><td>77.0</td><td></td><td>6.48</td><td>6.98</td><td>2.35</td><td>1.09</td><td></td><td>11.30</td><td>12.03</td><td>7</td></t<>		<b>429088</b>	AA005186	77.0		6.48	6.98	2.35	1.09		11.30	12.03	7
764356         AA436142         8.25         44.30         3.18         1.16         22.31         1.49         3.68         0.31           768226         AA42400         0.06         0.31         5.18         1.16         22.31         1.49         3.68         0.43           303139         N90806         0.52         2.71         5.17         4.05         23.41         0.96         1.45         0.13           757165         AA443950         0.03         0.16         5.09         0.47         1.51         0.94         20.25         2.06           757165         AA443950         0.03         0.16         5.09         0.47         1.51         0.94         20.25         2.06           247103         N53959         0.05         0.22         4.82         3.70         1.71         1.84         0.22         2.88           2247103         N53959         0.05         0.22         4.82         3.75         1.45         2.24         3.15         2.0           1 7550         AA438119         0.04         0.21         4.73         0.22         1.45         2.24         3.15         2.0           1 7555         H10881         0.10	2	28 24 28 28 28 28 28 28 28 28 28 28 28 28 28	NZ1Z33	3		¥ 7 ¥	2.04	2.77	1.1	7.77		4	8
768226         AA42490         0.06         0.31         5.10         4.05         23.41         0.96         1.45         0.13           303139         N90806         0.52         2.71         5.17         4.05         23.41         0.96         1.45         0.13           767165         AA443860         0.03         0.16         5.09         0.47         1.51         0.94         20.25         2.06           865235         AA43816         0.03         0.16         5.09         0.47         1.51         0.94         20.25         2.06           247103         N53859         0.05         0.22         4.82         3.70         1.71         1.84         0.22         2.88           224376         AV2288         2.94         14.16         4.81         1.46         2.58         4.82         2.24         3.15         2.40           758405         AA482119         0.04         0.21         4.73         0.22         1.45         2.24         3.15         2.40           47355         H10881         0.10         0.48         4.85         1.45         1.450         1.23         1.41         1.47           1377898         AA777088	=	754358	AA438142	8.25	28. Y	7	1 16	22.31	1.49	3.68	18.0 1	2 2	•
303139         N90806         0.52         2.71         5.17         4.05         1.51         0.94         20.25         2.06           757165         AA443850         0.03         0.16         5.09         0.47         1.51         0.94         20.25         2.06           86525         AA43812         18.25         88.08         4.88         7.86         1.443         2.52         1.47         0.61           247103         N53859         0.05         0.22         4.82         3.70         1.71         1.84         0.22         2.88           234376         N28289         2.94         14.16         4.81         1.46         2.58         4.82         2.24         3.15         2.40           234376         N28288         2.94         14.18         4.73         0.22         1.45         2.24         3.15         2.40           47355         H10881         0.04         0.21         4.73         0.22         1.45         1.29         2.30           47355         H10881         0.10         0.48         4.38         5.72         2.44         11.89         1.29         2.30           45072         AM5072         AM508275 <t< td=""><td>Ω</td><td>768228</td><td>AA424900</td><td>90.0</td><td>0.31</td><td>5.16 5.16</td><td>20.5</td><td>23.41</td><td>0.96</td><td>1,45</td><td>0.13</td><td>98.0</td><td>• •</td></t<>	Ω	768228	AA424900	90.0	0.31	5.16 5.16	20.5	23.41	0.96	1,45	0.13	98.0	• •
757165         AA443860         0.03         0.16         5.09         0.47         1.71         1.84         0.22         2.88           86525         AA13812         18.25         88.08         4.88         7.86         14.43         2.52         1.47         0.61           247103         NS3859         0.05         0.22         4.82         3.70         1.71         1.84         0.22         2.88           2347103         NS3859         0.05         0.22         4.82         3.70         1.71         1.84         0.22         2.80           234376         NS2858         2.94         14.16         4.31         1.45         2.24         3.15         2.40           756405         AA482119         0.04         0.21         4.73         0.22         14.35         0.22         9.85           47355         H10881         0.10         0.48         4.85         14.30         0.10         0.55         0.22         9.85           47355         H10881         0.10         0.48         4.85         4.38         5.72         2.44         11.65         1.29         2.30           74507         AM5072         AM52875         0.45         <	2	300139	N90806	0.52	2.71	5.17	SD: •	7	9.	20.25	2.06	5.30	•
666235         AA136126         18.25         68.08         4.86         7.86         1.71         1.84         0.22         2.86           247103         N53959         0.05         0.22         4.82         3.70         1.71         1.84         0.22         2.86           234376         N53959         0.05         0.22         4.82         3.75         14.34         2.24         3.15         2.40           756405         AA482119         0.04         0.21         4.73         0.22         14.5         2.24         3.15         2.40           47355         H10881         0.10         0.48         4.85         14.30         0.10         0.55         0.22         8.85           1 377898         AA777088         3.49         15.28         4.38         5.72         2.44         11.69         1.47         1.47           745072         AA626275         0.45         1.85         4.32         6.41         14.50         1.23         1.41         1.47	. 3	757165	AA443950	0.03	0.18	5.09	0.47	5.5	2 52	1.47	. 0.61	2.40	~
247103         NS3859         0.05         0.22         4.82         3.70         1,71         1.73         1.34           234376         NS3859         0.05         0.02         1.46         2.58         4.82         2.62         14.34           234376         NS3858         2.94         14.16         4.81         1.45         2.24         3.15         2.40           47355         H10881         0.04         0.21         4.73         0.10         0.55         0.22         8.85           47355         H10881         0.10         0.48         4.85         14.30         0.10         0.45         1.29         2.30           377898         AA777088         3.49         15.28         4.32         6.41         14.50         1.23         1.41         1.47           745072         AA626275         0.45         1.85         4.32         6.41         14.50         1.23         1.41         1.47	<u> </u>	KRK235	AA138125	18.25	89.08	4.88	7.86	24.4	1 7	073	2.88	18.58	~
234376         N28288         2.94         14.16         4.81         1.46         2.58         4.02         2.40           25405         AA482119         0.04         0.21         4.73         0.22         14.5         2.24         3.15         2.40           47355         H10881         0.04         0.21         4.73         0.10         0.55         0.22         8.85           47355         H10881         0.10         0.48         4.38         5.72         2.44         11.68         1.29         2.30           377898         AA777088         3.49         15.28         4.32         6.41         14.50         1.23         1.41         1.47           745072         AA628275         0.45         1.85         4.32         6.41         14.50         1.23         1.41         1.47	2 9	247403	N43959	0.05	0.22	4.82	3.70	U.T.		2.82	14,34	2.88	~
4735     0.22     1.45     2.24       758405     AA462119     0.04     0.21     4.73     0.22     9.85       47355     H10981     0.10     0.48     4.85     14.30     0.10     0.55     0.22     9.85       47355     H10981     0.10     0.48     4.85     14.30     2.44     11.68     1.29     2.30       745072     AA626275     0.45     1.85     4.32     6.41     14.50     1.23     1.41     1.47	2 !	201112	880001	294	14.18	4.81	1.46	2.58	7.07	3.45	2.40	18.90	~
756403 AAA02119 0.00 0.48 4.85 14.30 0.10 0.55 0.22 0.22 0.22 0.22 0.22 0.22 0.2	ă	010407		2	0.24	4.73	0.22	1.45	577	, ,	9.85	2.80	7
4735 H1080 3.19 15.28 4.38 5.72 2.14 11.08 1.29 1.47 1.47 1.47 1.45072 AA628275 0.45 1.85 4.32 6.41 1.450 1.23 1.41 1.47	<b>8</b>	736403	_	5	0.48	4.85	14.30	0.10	0.55	7 7	2.30	2.82	2
377888 AA777088 3.49 1.85 4.32 6.41 14.50 1.23 1.41 1.23 1.41 1.23 1.41 1.23 1.41 1.41 1.42 1.42 1.42 1.43 1.41 1.42 1.42 1.42 1.43 1.43 1.43 1.43 1.43 1.43 1.43 1.43	æ	47355		2 9	45.28	4.38	5.72	2.44	11.68	97.1	17.	0.83	~
745072 AA626275 0.45 1.83 7.02	2	377898	-	R .	3.5	133	6.41	14.50	1.23	Į.	<u>.</u>		
	Ξ	745072	_	0.45	CR.	10:1		;					

Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

							7-1-1-10M194/A	Fold/MPM124/A	Fold(MPM122/A	FOLIMPHIZAJA FOLDIMPHIZZA FOLDIMPHIZZJA FOLDIMPHIZGIA # OFTUMOT	old(MPM125/A	# of Tumor
	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTurnon/A	Fold(AveTumo#Av Fold(MPM120/A Fold(MPM121/A Fold(AveTumo#AveBPH)	rote(mrmiting)	veBPH)	veBPH)	veBPH)	VeBPH)	Seidmis
	Clone ID	Number				•	1		1.48	90'0	14.64	- 2
	707R3	TRIORY	0.16	0.71	4.32	1.17	7.49	70'1		0.83	4.17	7
<b>1</b>	3 6	100001	2	110	4.20	2.13	15.88	0.30	1.78	95	11.0	8
200	7747	A400100	3 6	98.0	4.17	16.96	1.24	1.87	20	900	1 42	. 7
3/6	00000	0077		5	4.16	2.17	2.93	3.20	13.79	ŧ ;	148	
375	1606315	SALLBAN			11.7	11.10	10.86	0.50	0.52	170	2 6	, ,
376	1386018	AA855158	6.6	9 6		1.22	10.53	5.68	2.67	1.88	98	• •
377	283335	N51441	0.18 8	0.72	00.4	1 G	, &	0.83	0.57	0.82	10.57	7
378	284931	N71861	0.02	0.07	3.99	2.68	900	43	1.64	0.36	15.08	~
370	1031203	AA609982	0.03	0.1	3.96	1.17	0.50		4.17	0.30	16.49	64
380	758360	AA404290	0.03	0.13	3.85	0.30	0.30	2 5	2.97	1.40	10.58	7
8	178569	H49053	0.05	0.20	3.85	3.16	3.00	4.58	167	2.16	12.53	~
382	184477	R54855	10.1	3.85	3.92	2.32	1.75	3.10		1.03	0.58	~
5	1048998	AA778638	0.14	0.57	3.91	10.73	5.96	7.90 7.90	1 6	3.45	1.09	8
7	129589	R16541	4.0	1.68	3.79	2.77	13.43	80°-	5	1.71	9.27	7
28.	1055636	AA620927	0.10	0.72	3.75	3.32	2.92	80:7		8.08	2.5	~
3 5	44.699	AAB82274	10.27	38.51	3.75	0.61	2.58	2.98		5.65	6.87	7
3 5	50892	H17626	0,31	1.18	3.74	1.87	2.97	2.88	177	12.0	1.0	~
3 8	2000	AAAABBBA	101	9.0	3.74	2.76	5.27	<del>.</del>	);;;	62.0	10.81	7
8	20404	T0084E	600	0.11	3.69	0.65	8.21	7.	<u> </u>		0.85	8
3	0/1771	CIDORI	8 6	2.06	3.69	6.45	12.29	22.	<b>3</b> .0	300	66	2
8	246808	4125CN	3 6	2,7	80.5	2.33	6.78	2.63	2.98	B. C.		
30	101888	H40350	5	47.		2 00	9.56	2.16	0.92	0.72	3	
385	155072	R71393	0.30	1.07	3.07	 ar.	2.71	2.34	2:82	7.51	3,33	<b>N</b>
88	190325	H29897	0.70	2.51	3.58	6.70		2.17	7.62	1.62	6.75	~
ğ	502438	A134862	1.10	38	3.58	1.28	• °	. F	1.03	10.18	1.08	~
385	283889	N69675	0.0	0.0	3.57	6.85	9 6	87.0	10.40	1.10	2.70	7
398	1488752	AA912071	0.70	2.48	3.57	1.94	9.0	1 88	1.98	1.85	2.87	~
397	510138	AA053239	0.03	0.10	3.52	6.17	0.37	3 6	7.18	5.88	2.39	7
388	878284	AA670280	2.85	9.83	3.45	1.42	0.10 0.00		2,62	2.54	1.99	8
388	321470	W32303	1.07	3.68	3,45	3.70	9 :	Ę	1.12	7.17	1.71	~
<b>\$</b>	267868	N62332	124	4.26	3.44	2.88	69.4	5.1°	7.38	2.52	1.53	~
5	258300	N30680	5.10	17.12	3.36	2.43	3,55 0 1	2.70	2.20	2.53	6.91	~
405	1048646	AA608893	0.82	3.08	3.35	1.86	2.72	1000	8 82	1.99	1.86	~
\$	289025	N59626	99.0	1.82	3.35	2.20	87 °	- e	5 6	1.50	1,49	~
\$	291708	N73480	0.53	1.77	3,33	4.14	T.8.1	4	2.61	1.62	4,45	8
405	1031568	AA608304	80.0	0.25	3.30	6.38	2.87					
							12					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

					of Tumol				Alcondary.	Fold(MPM123/A	Fold(MPM126/A	# of Tumo!
		400			Fold/AveTumor/Av	Fold(MPM1201A	Fold(MPM121/A	Fold(MPM124A	veBPH)	veBPH)	veBPH)	
	IMAGE	Gell Dallin	AveBPH	AveTumof	(Hdae	veBPH)	VeBPH)			9, 1	1,81	2
_	Clone ID				<u>:</u>		30.6	2.77	5.08	8.5.4 1.00	184	7
			,	170	3.28	7.55	9 9	2.54	4.55	5.37		C
<b>6</b> 04	383676	AA679116	0.13	F	1.28	2.57	2.10	7 7	2.78	0.15	0.69	4
	240493	W31074	6.22	20.37	07:0	585	6.18	2.87	2 6	4.14	0.35	7
Ì		13671	0.0	0.21	3.27	9 9	0.51	0.73	12.99		8.83	7
<b>4</b> 80	2 2		9	0.61	3.24	0.73	,	1.65	2.27	7	2	74
<b>6</b>	210955	H81517	<u>.</u>		3.23	2.04	<b>*</b> 1.7		8.52	1.58	n i	
410	285741	W02483	0.35	5.13		1.38	0.93	A0'L	88 6	1.48	10.68	7
;	708848	AA480115	9.12	29.44	3.63	55	1.21	0.62	8 8	3.85	2.98	~
;	67677	H05818	90.0	0.21	3.23	200	2.85	4.57	BO:7	3 02	1.53	~
1	120051	R62928	<b>5</b> .0	1.43	373	, g	1.39	2.00	06.01	7.72	2.25	7
2	784843	AA677023	0.83	2.67	322	9 6	1.60	<del>1</del> .	5.55	126	0.62	n
=	200	LISKIER	0.22	0.71	321	60.0	9.86	0.85	1.50		1.53	2
5 3	48313	W74462	1.22	3.89	3.17	08.7	4.47	2.58	6.53	2 E	0.98	2
5	310045	U00788	1.11	3.52	3.17	64.7	85.	1.99	4.43	70.7	2.01	84
\$	263630	POTERIL	2	2.28	3.15	7.49	39	1.85	7.85	<b>4</b>	90	2
418	376783	AA046883		69.6	3.13	0.71	<b>8</b> 9	2.75	1.53	0.65	8.5	. •
<del>+</del>	283617	N52875	1.18	90.5	3.53	5.71	7.10	6.73	3.58	6.73	1.25	4 (
420	1684274	A1003028	1.45	¥.	2 7	2.46	2.18	7. 7.		0.17	9.50	<b>8</b>
\$	784212	AA446864	1.91	5.95	- 3	0.98	3.88	<del>,</del>	S. 5	0.37	2.48	7
;	41214	R56898	0.03	0.11	11.8	7.63	1.12	<b>4</b> .50	7:37	33	2.00	<b>7</b>
ָּבְּיבָּיבָּיבָּיבָּיבָּיבָיבָיבָיבָיבָיבָיבָיבָיבָיבָיבָיבָיבָי	RAFAR		0.03	0.08	3.10	Ę	1.07	3.65	20.00	707	4.03	7
3	440043		7.60	23.51	3.09	. 60	2.94	2.83	2.19	77	0.99	7
•	200000		0.52	1.59	3.09	55.3	0.30	0.86	3.29	= 8 5	8.48	7
624	900700		0.0	0.20	3.08	60.21	1.48	1,52	1.66	3 8	091	2
426	20/5			1.61	3.08	87.1	2 7 3	1.58	1.33	00.0	4.2A	7
427			•	5.87	3.07	5.36	2.6	5.26	5.17	1.89	720	7
428	364000			3.80	3.07	2.60	2.87	3.48	1.95	1.40	4.82	7
2				0.13	3.07		2.85	2.36	1.7	15.4	5.35	8
3 3				90.08	3.07	84.7 84.7	2.60	4.93	1.72	2.4	4.86	8
<b>7</b>				1.85	3.07	2.6	1.81	2.18	2.33		1.15	7
3				5 1.07	3.04	2.80	88.6	1.88	2.92	2.40	7	7
<del>1</del>				2 0.37	9. 9.	3.96	5.47	4.57	1.88	2.74		7
\$				9 0.48	3.04	2.44	F 24	1.48	2.88	3.18	10.	2
435					3.0	2.94	77.5	1 10	9.90	3.38	5.	•
<b>98</b>	••				3.03	1.17	\$ <b>9</b>	77.	2.64	4.52	8.7	. ~
5					3 3.02	2.49	5 6	1.38	2.38	4.05	3.4	•
<b>\$</b>		•			3.01	2.74	70.7					
439	9 261580	70 H98694					5					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession Number	AveBPH.	AveTumor	Fold(AveTumor/A eBPH)	Fold(AveTumoriav Fold(MPM121/A Fold(MPM12L/A Fold(MPM122/A Fold(MPM122/A # of Tl «BPH) veBPH) veBPH) veBPH) sams	Fold(MPM121/A veBPH)	Fold[MPM124/A voBPH]	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(MPM126/A veBPH)	# of Tt.
2	357628		0.12	0.37	3.01	3.08	1.37	8.10	2.01	2.27	123	7
2	701766	AA292655	4.22	12,71	3.01	1.74	5.33	1.83	2.67	2.63	3.85	~
442	195487	R82163	1.07	3.20	3.00	2.24	4.24	4.46	1.57	2.77	2.74	7
45	480184	AA678920	40	1.32	2.89	1.42	2.73	1.25	1.78	6.89	4.80	8
\$	788239	AA478452	1.47	4.37	2.98	4.17	6.87	1.68	1.67	2.47	1.06	7
445	365813	AA025601	90.0	0.18	2.87	2.12	6.90	1.18	4.61	9.14	2.91	~
31	346889	W79834	7.57	22.52	2.87	2.25	2.78	1.65	2.96	4.30	3.91	~
#	143780	R76782	2.09	6.21	2.97	2.52	2.84	2.73	4.48	3.12	2.14	~
2	60530	H17919	0.35	<b>7</b> .	2.86	5.5	2.38	1.71	3.99	6.14	2.52	~
2	208082	H62563	4,23	12.48	2.86	2.28	6.36	2,48	4.67	0.81	1.18	7
\$	480525	AA128488	0.07	0.20	2.85	0.16	1.59	10.45	1.7	0.15	3.62	8
. 451	264157	N20602	0.08	0.28	2.85	4.52	5.80	2.94	2.25	0.12	2.05	7
452	133440	R27450	1.72	5.05	2.94	2.28	2.49	1.50	. 5.90	3.21	2.23	~
453	416889	W86832	0.52	1.62	2.82	1.81	5.15	4:77	1.53	1.35	2.83	~
\$	271078	N29918	2.08	6.09	2.82	4.05	4.90	2.55	2.04	1.62	2.38	7
455	452320	AA704777	1.07	3.13	2.82	2.22	2.90	2.18	3.07	2.67	4.48	~
456	814798	AA455235	25.12	73.30	2.82	3.83	7.12	1.86	2.07	1.26	1.56	~
457	417908	W90520	5.87	17.42	2.92	2.62	6.05	2.91	4.81	0.62	0.47	7
458	745283	AA625587	5.50	16.00	2.91	2.17	1.35	1.50	2.45	5.86	4.13	7
459	50772	H16803	10.55	30.44	2.89	3.09	5.75	2.12	2.83	1.37	2.15 -	7
460	51608	H18938	0.68	1.98	2.88	2.55	3.73	3.69	2.18	2.15	2.70	~
461	204684	H57135	0.83	2.38	2.88	1.27	2.47	2.58	4.88	4.06	2.03	7
462	731002	AA16864	1.11	3.20	2.87	2.09	3.10	4.25	2.58	2.78	2.48	7
<b>4</b> 83	124891	R06119	0.85	2.43	2.87	2.66	3.84	3.59	2.24	2.73	2.16	~
\$	428138	AA002007	97.0	1.58	2.87	2.46	3.04	2.60	5.16	2.82	<b>:</b>	7
485	45587	H07934	69.	4.82	2.85	2.11	5.17	3.34	2.83	1.61	2.04	7
466	731254	AA420989	90.0	0.13	2.84	2.74	1.81	3.89	1.81	27	4.51	~
487	48055	H08850	0.74	2.09	2.84	0.41	0.65	0.73	4.40	10.07	0.75	N
468	626385	AA189113	29.81	<b>8</b> .3	2.83	2.15	2.80	3.18	6.89	1.20	0.76	7
469	276450	N34824	0.47	1.32	2.62	6.04	3.10	1.85	1,41	2.48	2.16	84
410	460435	AA677581	3.97	11.17	2.82	2.78	3.27	2.11	2.34	4.55	1.85	~
41	41825	R54109	0.07	0.21	2.82	0.83	4,13	2.29	2.38	0.14	7.15	8
17	682058	AA256458	1.86	5.21	2.80	4.23	4.71	2.40	2.75	0.67	2.03	2
£	754408	AA438187	0.03	60.0	2.78	5.62	0.31	1.97	2.18	4.95	1.67	~
						4	-					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank	AVARDH	AveTumor	Fold(AveTumoriav Fold(MPM120IA Fold(MPM121IA Fold(MPM122IA Fold(MPM123IA Fold(MPM126IA * of Tumorian * of Samples * of Sam	Fold(MPM120/A	Fold(MPM121/A vèBPH)	Fold(MPM124/A F	old(MPM122/A	Fold(MPM123/A veBPH)	Fold(MPM126// veBPH)	Samples
	Clone ID	Accession			(HA80		•		4 4	6,00	2.88	7
			,	600	2 78	2.12	2.47	7.7	2 6	į	2.89	2
<b>11</b>	4994	H28215	13.79	30.3	F 6	1,48	3.14	4.59	2.82		127	74
475	291222	N72196	0.08	0.23	7:10	. 6	2.81	2.83	0.95	7.30 7.30		
478	789857	AA430367	1.25	3.45	2.78	07.5	2.80	89.4	2.49	2.42	9.3 9.1	۷ (
Ę	195875	R92216	0.58	<del>1</del>	2.78	88.0	2 5	0.88	4.74	2.38	ž Ž	<b>7</b>
Ę	731048	AA421275	0.02	0.05	2.76	0.58	7.7	264	3.65	1.52	2.70	7
	01308K	AA484800	0.87	2.41	2.76	4.36	1.66	5 6	776	1.72	1.19	7
	202710	94744	0.17	0.48	2.78	1.66	5.83	3 3		2.28	6.97	7
480	897808	A4434730	: 8	¥	2.75	0.51	3.74	<b>7</b> .24		40.4	6.80	7
<b>\$</b>	288513	N62745	70.0	3 ;	2.75	2.08	1.72	3.12	1./4	9 9	2.78	7
482	327239	AA284307	0.50	ري د د د		2.22	4.10	3.71	2:22	g (	7	- 24
£83	23588	. R38369	1.07	2.96	5.73	52.	3,42	2.33	2.85	A/.L	i d	
\$	153817	R48833	1.27	3.49	2.75		¥ 33	2.24	4.29	1.74	98.7	
485	491435	AA150435	0.28	0.78	2.75	58:0		88.0	0.55	8.93	<b>4</b> .84	2
3 9	238782	HR1193	0.12	<b>7</b> .0	2.75	4,13	0.00	5 5	3.17	0.83	7.34	7
Ş ;	4070C7	001121	0.08	27	2.75	2.78	0.80	8 6	1 98	1.26	1.51	7
9	101			2.71	2.74	7.49	3.35	0.02	2 6	1.07	1.83	7
<b>\$</b>	786525	AA452113	5 6		2.74	2.57	4.19	₽. <del>.</del> 4	10.7		2.31	64
489	742082	AA405769	D. 13	g (		2.05	2.54	2.17	3.06	3 6	24	7
8	38821	R53428	12.95	35.28	2.7	75	4.72	4.82	2.58	0.83		
491	40338	R53072	2.38	8. 25.	2.73	, i	1 67	1.04	1.82	4.30	80.4	4 (
492	194638		10.30	28,05	2.72	177	5	3.46	4.87	1.74	4.77	7
167	149408		3.58	9.74	2.72	1.80	5	1 22	0.80	1.83	1.89	7
3			6.45	17.51	2.71	3.86	94.9	1 2	1.20	0.70	7.49	7
ţ ;	15.170B		0.71	<b>.</b>	2.71	2.51	¥. ;		2.48	3.61	2.51	8
C S	000000	•	1.30	3.51	2.70	1.67	3.06	90.7 0	80	2,28	3.88	7
\$ !	000074	•	0.52	1.40	2.70	<u>4</u> .	2.59	9.60	36.	4.19	2:32	2
497	65430	-	50.0	0.14	2.70	4.16	4.07	28.5	\$ 5°	1.89	1.66	2
<b>4</b>	PO 107		133	3.57	2.69	1.60	2.33	3 (		54.5	2.71	2
88				8.39	2.69	3.78	4.41	1.27	3 5	4.20	2.33	7
8			3 4	10.81	2.69	2.74	3.62	1.27	è .	£.	2.60	74
25	878281	•	,	4	2.68	1.36	3.04	2.45	1.82	6	0.82	8
802	19291		CR.41	<u> </u>	268	0.63	2.98	7.53	3.18	. 99.0	2	2
200	625924	AA188885	0.08	17.0		3.26	2.72	2.12	2.74	0.7		ור
Š	810328	1 AA464143	0.28	0.75	7.07	24.0	6.02	0.45	5.94	0.35	0.0	۷ (
505	50114	H16743	2.07	5.53	2.67	6.3	151	4.67	0.86	421	88.	7 (
8	743731	AA628357	0.28	0.76	2.67	28.7 7	1.0.1	125	3.28	¥.	2.28	Ν
507	-	AA707118	0.41	1.08	2.67	1,41						
•							15					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank	100		Fold(AveTumoriay Fold(MPM120)A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM125/A # of Tum	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	old(MPM125/A	# of Tun
	Clone (D	Number			eBPH)	veBPH)	veBPH)	VeBPH)	VeBPH)	(MAREA)		
8	381587	AA016980	0.88	2.34	2.65	2.25	2.02	1.7	1.78	3.89	4.19	64
809	283142	N51336	4.33	11.48	2.66	1.58	2.50	0.65	5.68	3.31	77	7
. 20	342721	W88291	0.58	1.53	2.65	1.63	2.24	3.17	6.41	1,43	1,03	~
512	530185	AA111969	0.31	0.83	2.65	4.40	3.73	2.69	2.49	0.20	2.38	8
512	395609	AA757588	0.03	0.0	2.65	7.24	3.66	5.09	1.38	0.84	0.68	<b>84</b> .
513	898318	AA598828	2.48	6.52	2.65	3.03	4.78.	1.15	2.49	<del>1</del> .	3.00	7
514	241787	H91724	0.16	0.42	2.65	1.72	3.98	3.91	1.88	1.88	2.52	8
515	1636108	AID15679	2.30	6.08	2.65	2.96	2.48	3.78	1.84	3.28	5.7	7
516	201172	R98487	0.87	1.78	2.65	1.56	4.18	2.12	1.90	2.78	3.37	~
517	229290	H63865	0.70	1.86	26.	1.33	2.40	197	1.57	4.53	8£. <del>4</del>	7
8,10	342164	WB3778	9	0.10	25.	0.60	0.28	5.08	1.19	0.28	8.48	~
510	48384	H15855	0.51	1.35	28.	2.46	4.41	2.08	2.88	3.04	86.0	~
23	85320	771889	6.68	17.60	25	1.58	1.83	4.01	2.90	3.08	.230	8
521	115143	188708	5.38	14.12	2.63	3.65	6.57	1,85	1.85	0.94	1.14	~
23	31960	R41994	0.74	1.96	2.63	1.11	1.08	1.83	1.88	5.62	4.28	~
523	362402	AA018449	8.88	25.98	2.63	2.80	4.67	2.27	3.51	1.83	<b>5</b> .	7
624	283049	N20048	0.35	0.91	2.63	1.09	4.25	1.58	7.15	0.95	77.0	7
625	31299	R42948	0.73	1.92	2.63	1.79	3.06	2.41	3.54	2.21	2,78	7
528	62112	141077	3.82	10.27	2.62	3.22	2.98	1,49	3.17	2.47	2.40	7
527	49728	H29198	0.70	1.83	2.61	2.66	2.18	3.18	1.50	2.17	3.98	7
828	205723	N66925	2.82	7.37	2.61	2.25	2.74	3.72	1.30	2.40	325	7
228	447715	AA702781	14,05	38,83	2.61	2.78	3.25	1.62	4.68	2.16	<del>2</del>	7
8	828070	AA588779	0.82	2.40	2.61	1.58	2.75	1.77	2.53	3.64	3.38	7
2	60540	140531	1.12	2.91	2.60	2.02	2.15	2,44	2.01	3.34	3.68	~
g	288899	N82728	3.95	10.29	2.60	2.27	3.83	1.45	3.80	2.10	2.08	~
83	280348	N62283	1.53	3.89	2.80	1.83	3.19	1.53	4.80	1.98	2.40	~
ğ	51865	H23187	0.05	0.12	2.59	2.52	1.93	1.70	3.83	3.45	2.14	~
55	815124	AA481491	1.93	5.02	2.59	1.28	1.68	2.69	3.39	4.11	2,42	~
8	428298	AA004944	09.0	1.56	2.59	1.85	2.54	2.85	1.58	323	3.47	~
537	51607	H18427	0.15	0.40	2.58	2.50	5.85	3.09	0.82	2.24	0.98	~
8	283461	N50847	90.0	0.14	2.58	1.74	2.31	0.18	0.37	7.68	322	7
82	283190	N51357	0.07	0.19	2.58	1.89	5.87	2.69	3.98	0.14	0.82	64
540	74511	T59024	10.71	27.80	2.58	3.63	2.79	2.02	4.23	1.30	1,49	8
3	300051	N78927	1.59	4.10	2.57	3.34	0.28	0.12	0.68	10.78	0.24	7
						=	18					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

AVEBPH 1.38 1.79 0.07 8.33 0.37 0.29	AveTumor	Fold(AveTumonAv Folution weBPH) veBPH) veBPH	veBPH)	veBPH)	veBPH)	(U.JOBA		•	•
								1,95	7
1.38 1.79 0.07 8.33 0.37 0.29				15.1	1.84	3.37	8/G	1.15	7
0.07 9.33 0.37 0.29	3.59	2.57	5 5 7	1.9.1	1,52	4.86	A. 4	1.13	7
0.07 8.33 0.37 0.29	<b>4</b> .60	2.67	77:	2.47	2.87	3.13	S 1	223	7
6.33 0.37 0.29	0.19	2.57	5 <del>.</del> .	1 70	3.19	1.45		1.05	7
0.37	23.95	2.57	£	253	2.58	2.31	7G.E	9	~
0.29	0.98	2.58	1.1	;	2.09	0.54	3.25		6
0.07	72.0	2.58	2.80	R. G	27.6	3.32	1.1	71.1	, «
0.0		2.56	4.68	2.29	7.7a	1.95	4.73	2.53	<b>,</b> (
	81.0	2 4	1.19	1.84	30.5		5.94	3.87	7
0.49	27.	5.00°2	1.72	1.10	1.28	<u> </u>	2.81	4.51	~
0.22	0.58	25.4	323	2.89	0.80	5 6	3.30	2.72	7
9.34	0.88	6.33	2 5B	3.35	0.81	8 · 7	75	3.60	7
3.70	9.38	5.54	22.0	2.86	1.99	2.48	1.48	1.59	2
1.10	2.78	2.54	. 25	3.69	1.78	4.20	89	2.38	7
2.03	5.15	2.54	3 4	2.61	3.14	1.88	20.7	99:	~
8.02	15.28	2.54	5	4.28	1,05	1.58	7 C	3.89	7
0.57	<del>1</del> .	2.54	<u> </u>	1.48	2.26	3.1	70.7	58.0	~
3.91	8.80	2.53	S .	3.90	1.93	1.64	Š.	9.38	2
AA443140 0.14	9.3	2.53	2.23	9 45 16 16	1.80	2.58	1./3	1 1	~
	2,01	2.53	3.12	3 6	121	3.56	1.90		~
7.	4.38	2.52	2.94	0.5	3.28	1.83	3.20	76.7	. ~
		2.51	2.69	1.78	3 6	3.11	3.36	2.05	
AA098593 3.24	2 6	2.51	1.7	1.90	70.2	2.23	1.78	1.57	7
0.50	1,26		3.86	4.71	5.0	95	0.86	<u>+</u>	7
1.39	3.50	. 63 c	3,52	6.33	1.90	95.	1.77	1.78	~
11.98	28.82	2.50 8.50	1,35	1.80	1.09	60.0	3.11	1.27	~
	68.89	9 9	2.80	0.09	4.82	7.50	2.62	1.91	7
AA156030 0.11	0.27	2.50	2.26	3.58	3.63	4.87	3.26	1.69	4
1.96		2.50	1.43	1.30	2.43	9 6	3.94	2.48	7
W73892 1.10	2.74		1.18	1.68	2.43	2 6	2.69	1.75	7
AA883299 0.94	2.35	8 ° ° °	1.32	2.39	3.08	3.07	3.20	2.96	7
AA043552 2.47	6.13	0.40 0.48	1.78	1.10	1.39	7 6	1,85	<u>.</u>	7
AA703048 0.74	.83	04.4	3.17	2.83	2.41	9.78	3.01	1.05	~
AA489639 0.69	<u>r.</u>	0 7	101	2.44	2.81		1.93	0.78	7
	5.1 5.2	7 C	2.13	423	3.13	1.0.4	128	¥.	~
H17921 0.38	0.83		2.30	3.42	77	5.5	1		
N86132 8.84	24.32	ř.	ı	11					
~~66		1.36 1.10 0.94 0.74 0.68 0.38	1.96 4.09 1.10 2.74 0.94 2.35 2.47 6.13 0.74 1.83 0.69 1.71 0.48 1.18 0.36 0.83 9.84 24.32	1.96 4.59 1.10 2.74 2.50 0.84 2.35 2.49 0.74 1.83 2.48 0.69 1.71 2.48 0.48 1.18 2.48 0.38 0.93 2.47 8.84 24.32 2.47	1.96 4.09 1.43 1.10 2.74 2.50 1.48 0.84 2.35 2.49 1.32 2.47 8.13 2.48 1.78 0.74 1.83 2.48 1.78 0.48 1.71 2.48 3.17 0.48 1.18 2.48 1.01 0.48 24.32 2.47 2.30	1.96     4.08     1.43     1.30       1.10     2.74     2.50     1.43     1.30       0.84     2.35     2.49     1.18     1.68       0.74     6.13     2.48     1.78     1.10       0.74     1.83     2.48     1.78     1.10       0.49     1.71     2.48     1.01     2.44       0.48     1.18     2.48     1.01     2.44       0.38     0.93     2.47     2.13     4.23       9.84     24.32     2.47     2.30     3.42       17     17	1.96     4.09     2.50     1.43     1.30     2.43       1.10     2.74     2.50     1.18     1.68     2.43       0.84     2.35     2.49     1.18     1.32     2.39     3.08       2.47     6.13     2.48     1.78     1.10     1.39       0.74     1.83     2.48     3.17     2.83     2.41       0.49     1.71     2.48     1.01     2.44     2.81       0.38     0.93     2.47     2.13     4.23     3.13       0.39     2.47     2.30     3.42     2.22       8.84     24.32     2.47     2.30     3.42     2.22       17     17	1.96     4.08     4.08     2.45     3.23       1.10     2.74     2.50     1.48     1.68     2.43     3.23       0.94     2.35     2.49     1.18     1.32     2.39     3.08     3.87       2.47     6.13     2.48     1.78     1.10     1.39     4.47       0.74     1.63     2.48     3.17     2.83     2.41     3.29       0.46     1.71     2.48     3.17     2.44     2.81     4.55       0.46     1.18     2.48     1.01     2.44     2.81     4.55       0.36     0.33     2.47     2.13     4.23     3.13     2.87       0.36     0.33     2.47     2.30     3.42     2.22     3.78       8.84     24.32     2.47     2.30     3.42     2.22     3.78	1.96     4.08     4.09     2.43     3.23     3.94       1.10     2.74     2.50     1.43     1.30     2.43     3.23     3.94       0.84     2.35     2.49     1.18     1.32     2.39     3.08     3.67     2.56       2.47     6.13     2.48     1.78     1.10     1.39     4.47     3.20       0.74     1.63     2.48     3.17     2.83     2.41     3.29     1.85       0.89     1.71     2.48     3.17     2.44     2.81     4.55     3.01       0.74     1.18     2.48     1.01     2.44     2.81     4.55     3.01       0.48     1.18     2.48     1.01     2.44     2.81     4.55     3.01       0.38     0.93     2.47     2.13     4.23     3.13     2.87     1.28       0.38     2.47     2.30     3.42     2.22     3.76     1.28       0.84     2.43     2.47     2.30     3.42     2.22     3.78     1.28

Table 2-1:T3NO vs. BPH; Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession Number	AveBPH	AveTumor	Fold(AveTumon'Av Fold(MPM120/A Fold(MPM120/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumol eBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH)	Fold(MPM120/A veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A veBPH)	Fold(MPM1267/ veBPH)	A & of Tumor Samples
678	200104	H83952	0.69	1.70	2.47	88	6.03	0.73	2.48	2.35	1.00	
211	308466	N95485	7.9	19.83	2.47	2.01	3.49	2.05	4.93	1.49	0.85	<b>7</b>
578	78584B	AA449120	9.03	0.07	2.48	3.20	5.89	1.70	2.59	0.85	0.35	8
579	-	H52448	0.15	0.37	2.46	1.18	1.24	4.12	5.65	0.95	1.63	7
989	288570	N82788	6.23	15.34	2.46	1.31	3.10	3.00	3.31	2.60	1.48	7
158	67069	170438	0.62	1.53	2.48	7.88	3.28	2.	96.0	0.82	0.71	8
283	25061	R38944	14.58	33.84	2.46	1.10	3.69	3.77	1.95	2.40	1.88	8
88	566501	AA151917.	2.97	7.28	2.45	2.15	1.50	2.68	1.02	3.10	4.28	8
282	745206	AA626864	0.53	1.30	2,45	1.24	2.32	2.05	2.52	3.50	3.07	7
288	785890	AA448362	6.28	15.39	2.45	3.01	2.79	1.45	1.12	2.10	4.21	2
586	358048	W84591	1970	1.48	2,45	3.60	6.39	1.82	1.21	1.17	0.50	8
587	563860	AA101173	0.03	90.0	2,45	3.25	74.0	1.58	0.77	1.51	7.13	7
588	795407	AA453518	0.13	0.32	2,44	0.33	0.97	1.47	5.45	0.62	5.82	7
589	1480895	AA890098	5.95	14.53	2,44	2.08	2.15	1.42	2.58	3.31	3.12	81
280	753162	AA400457	2.38	3.8 48.	2,44	3.25	3.62	0.79	1.82	2.27	2.80	7
691	712489	AA278382	2.63	6.41	2,44	1.50	3.82	2.40	3.47	1.58	1.88	7
592	489858	AA099369	9.0	0.10	777	4.08	1.37	2.01	1.00	0.24	6.92	~
583	110503	T88886	09.0	1.45	2.43	0.73	1.24	0.76	0.82	5.55	5.38	8
265	197485	H52048	0.36	0.85	2.43	1.53	2.49	3.8	0.91	4.24	1.47	7
585	429729	AA011685	1.61	3.91	2.42	2.40	4.68	1.51	3.19	1.84	0.93	8
286	731371	A4421047	0.03	0.07	2.42	0.99	3.84	2.29	2.95	0.33	4.12	8
597	782585	AA447527	0.48	1.17	2.42	3.45	1.49	5,47	1.78	1,4	0.90	7
588	811010	AA485385	90.0	0.14	2.42	1.99	2.41	3.45	1.97	0.58	4.13	8
288	743445	AA609365	3.21	7,78	2.42	2.19	3.40	1.76	1.31	2.53	3.33	2
8	307328	N93470	0.03	0.0	2.42	2.00	3.80	0.67	1,74	2.14	1,15	7
601	297439	W03887	24.42	59.00	2.42	2.70	4.38	0.99	4.78	1.41	0.25	8
. 209	245320	N53456	1.20	2.89	2.41	4.58	3.25	2.53	2.71	0.68	0.75	8
8	140057	R65963	0.76	1.82	2.41	1.19	2.77	2.50	3.00	1.88	3.13	8
<b>\$</b>	433294	AA699707	1.60	3.85	2.41	1.23	3.14	1.87	124	4.06	2.83	74
803	812227	AA455369	4.47	10.79	2.41	3.67	5.67	2.04	1.25	0.84	0.08	7
808	1580977	AA875010	1.07	2.57	2.41	2.29	2.67	3.01	3.69	1.52	1.27	8
607	283379	N52787	9. 2.	0.10	2.41	4.36	4.19	202	0.77	0.87	2.25	8
808	85195	171380	0.02	0.05	2.41	4.76	0.45	0.28	0.45	8.08	0.45	8
8	813310	AA455164	3.54	8.52	2.40	2.78	4.07	1.41	2.17	3.03	0.99	7
						•						

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		3 5 6			A Part of the Part	Fold(MPM120)A Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumo	A Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	semples
	IMAGE	Accession	AveBPH	AveTumor	FOIG(AVE) UIIIO	VeBPH) VeBPH)	veBPH)	(MARA)			•
	Clone 10	Number			<u>:</u>	80 6	3.15	5.03	1.94	0.69	7 (
6.5	727157	H85345	20.24	48.65	2.40		181	1.86	4.37	1.35	2
2 5	48008	AA056534	0.42	1.02	2.40		£ \$	2.47	3.18	4.01	7
- ¢	445677B	AA883314	1.81	4.35	2.40		20°C	3.84	3.19	1.14	7
710	1486540	A4976714	5.57	13.37	2.40		60.	2.54	1,47	1.39	8
5 4 A	197787	R93727	1.09	2.61	2.40		6.5	1.07	1.12	1.93	~
. E	731272	AA420993	3.88	9.29	2.40	5.00	17	5.7	1.61	2.01	7
2 5	501431	AA115300	1.00	2.30	2.39		03.50	2.18	4.08	1.48	<b>C4</b>
. 5	415583	W04318	0.76	1.81	2.39		4.55	5.28	0.90	<b>1</b> .8	7
5 6	388380	AA732832	0.10	0.24	2.39	•	40.	7	1.37	0.52	<b>~</b>
5 6	787991	AA418828	6.24	14.88	2.39		2.88	0.88	0.70	1.47	7
	40204	B81798	18.89	44.60	2.39		2 48	3.03	2.42	. 1.88	~
8 8	10R44	AA005386	1.75	4.16	2.38		2 5	2.49	0.50	6.73	7
3 8	272478	N35907	11.37	27.10	2.38		ט ער יי	¥.	2.81	1.24	7
7 6	127083	R07891	1.23	2.83	2.38		CO:-	4.28	1.43	0.91	2
3 6	704168	44456135	12.87	30.59	2.38			5.01	1.77	1.15	~
928 828	795827	AA481508	0.03	0.07	2.37		23.5	3.14	1.72	1.48	~
3 6	1802284		1.50	3.56	2.37		378	3.39	99.0	2.73	7
3 6	487859		0.20	0.47	2.37	7.7 2.2 2.3	1 87	3.81	2.50	<del>.</del> 08	~
	1460391	AA883812	0.78	1.84	2.37		3.22	1.78	4.47	1.74	~
82	22700	R43588	1.08	2.58	2.37		231	1.16	1.43	0.93	7 (
8	432581	AA699390	7.05	16.64	2.38		<b>1</b>	0.80	1.70	5.24	7
831	757222	AA496149		6.63	2.36		3.02	4.59	1.37	2.78	<b>178</b>
2	244898	N54512	0.26	79'0	2.38		2.31	2.33	1.24	3.15	2 (
8	413300	AA772487	0.36	0.84	2.38		1.08	1.43	3.15	1.69	N) (
ş	285443	NB6393	1.60	3.77	2.38	3.65	1.91	2.15	0.83	<del>2</del>	<b>,</b> ,
838	625857	AA504783	<u>.</u>	3.61	2.36		1.89	4.01	1.75	0 <del>2</del> 0	•
8	264938	N21237	4.33	10.18	2.35		1.17	0.79	0.75	£ :	٧ .
637	43709	H05734	0.57	<u></u>	2.35	4.30	3.80	0.62	4.07	2.43	N (
83	41720	R54105	0.09	0.21	2.35		1.37	1.81	Ę.	0.72	~ •
g	299679	W05828	0.81	1.89	2.34		2.39	1.15	3.48	3.75	N (
3	731180	`	2.0	1.51	表 :		1.79	3.21	3.01	2.78	7
2	610264	AA463932	0.60	1.40	7.3		2.88	3.18	0.12	208	7
842	898055		_	0.12	7.	3,48	0.74	17.0	0.91	0.97	7
<b>3</b>	1240220	AA788970	4.63	10.81	2.33	9					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumoríAv Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor Arbhi verphi verphi verph) verph) verph	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124A	Fold(MPM122/A veBPH)	Fold(MPM123/A   veBPH)	Fold(MPM126/A veBPH)	s of Tumor Samples
		Number			:				<b>4</b>	730	1.78	~
3	242037	H93328	0.38	0.89	2.33	3.07	2.13	0.93	3 6	1 8		
645	470914	AA032084	13.04	30.41	2.33	2.36	3.53	7.7	2.33	3	8	<b>u</b> (
2	38260	R49462	0.70	2	2.33	1.90	0.47	0.87	0.59	6.53	4.59	~
643	50250	H17800	. 0.81	1.89	2.32	3.30	3.08	2.73	2.02	1.50	1.32	~
3	7/32/7	AA400128	1.85	007	2.32	1.78	3.31	2.05	2.70	3.18	0.82	~
3	290841	N71987	8.0	10.65	2.32	3.15	3.16	0.87	2.32	1.70	2.7	~
Ş	814505	AAABOOGA	2	13.75	2.32	7.7	2.10	1.47	3.01	4.67	1.50	~
3 5	240768	HR0245	2.74	333	2.32	1,73	3.33	3.46	2.02	1.79	1.58	64
<b>3</b>	3489	BALLA7	2.35	279	2.31	1.25	. <del>1</del> .	1.59	3.78	2.65	3.37	~
٤	412A3	22827	80	2	2.35	4.10	1.12	1.78	1.85	1.37	3.63	~
. 2	81000	AA484572	1.10	2.54	2.30	1,35	0.96	5.47	1.37	3.40	1.28	۲۰
, y	705831	AAARIKOR	70 0	617	2.30	1.39	2.39	3.59	5.16	0.48	0.83	~
3 5	745138	AARORGB	12.78	29.36	2.30	3.25	2.34	3.04	2.67	1.74	0.77	7
\$	489076	AA057195	0.02	50.0	2.30	1.88	3.30	2.10	5.47	0.55	0.48	~
5	344108	W73784	3.42	7.83	2.28	4.00	3.17	2.58	1.61	0.98	1.43	7
8	207.727	A 4 2 0 2 8 1 0	12.78	29.26	2.29	2.09	0.89	2.24	3.41	3.56	1,55	~
8	41541	R52789	77	25.	2.29	2.18	3.57	3.59	1.28	1.48	1.60	· 7
8	785353	AA453487	4.31	9.88	2.29	1.68	4.15	1.50	4.58	0.88	0.94	~
682	6400	R52088	0.31	0.71	2.28	2.41	0.86	1.19	0.31	4.68	<b>4</b> ,28	~
8	48330	H14949	0.17	88.0	2.28	1.85	3.18	2.62	2.50	3.07	0.48	~
\$	769600	AA425900	0.09	0.20	2.28	3.18	0.12	2.08	2.75	5.09	0.47	~
665	112885	T87130	2.81	5.85	2.28	1.59	3.50	3.29	1.78	1.99	1.51	8
ş	362705	4401883	90.0	0.13	228	2.04	6.08	0.18	3.14	0.18	2.06	8
<b>§</b>	184207	AA708A7R	0.48	50.	227	1,47	3.59	1.54	1.86	3.17	1.91	~
88	28568	R39111	7.0	1.68	2.27	3.15	0.55	0.70	2.64	1.53	5.05	8
99	199228	R85811	0.58	1.27	127	1.69	3.64	3.07	1.83	1.63	1.65	
970	41070	R58100	0.03	0.07	227	1.48	2.18	0.33	1.18	4.15	4.27	~
67	51103	H19217	0.37	0.83	227	0.74	4.79	¥.	1.55	3.13	z.	7
278	346552	W74377	0.02	20.0	227	3.52	2.57	3.03	2.05	0.61	1.81	~
E	624785	AA181995	4.34	9.82	2,28	3.02	3.29	1.52	2.48	1.11	1.51	~
674	626716	AA191548	17.62	39.85	228	3.68	3.59	0.94	2.27	1.42	1.67	~
675	785780	AA449686	0.02	9.0	2.28	3.56	3.49	2.71	2.58	0.71	0.53	~
678	345145	W72231	3.57	8.07	2.28	3.73	3.52	2.16	1.82	0.60	1.82	~
110	46620	H10038	15.87	36.82	2.28	0.96	1.62	1.52	1,48	3.82	4.13	~
							20					í

Table 2-1;T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	1000	Gen Bank	i		Fold(AveTumoriAv Fold(MPM120IA Fold(MPM124IA Fold(MPM124IA VeBPH) veBPH) veBPH) veBPH) Samples	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	veBPH)	veBPH)	V•BPH)	Samples
•		Accession	AVBER	Averamor	eBPH)	Veurn)			i	10.4	4.87	7
_		Number			•	78.0	0.21	2.33	LZ:0	,	2.78	
	811832	AA483827	90.0	0.11	2.25	ò	2.11	3.19	5.73	1.38	9	
		1340664	91.0	7.0	2.25	76.1		ş	1.38	1.62	7.70	• •
	48141	100 L		£ 8.7	224	3.20	3.06	<u> </u>	78.	0.37	0.38	7
	823891	AA489752	2.53	0.0	į ?	2.21	5.87	0.68	<b>.</b>	77.7	1,75	7
	754127	AA478827	3.54	7.92	3 5	02.4	<u>\$</u> .	98.0	3.17		4.63	7
,	4481809	AA883800	0.20	Ξ.	2,23	2 .	4 17	1.42	3.53	77.0		•
	2000	105140	0.32	0.70	22	1.86	- 6	76.	1.7	1.67	1.72	4 6
	200	01.00	2 2	48.25	22	3.16	50.5		3.48	0.79	1.35	7
	232912	H72643	7	30	2.21	1.97	3.06	60.7	2.08	1.72	0.28	7
	135789	R33353	r	3. 6	221	3.92	3.22	1.18	20.4	1.21	3.10	~
	795308	AA454149	0.10	67.0		2.46	1.00	3.23	2.25	H.	3.05	2
	448270	AA7777686	90.0	0.13	177	4 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	3.01	0.59	2.16		3.28	7
	898195	AA598587	1.7	3.7	777	2	1.24	1.02	5.17	6.0	000	7
	131073	R23241	4.58	10.05	2.20	5	3.15	3.31	2.10	1.88	9 6	
	77777	76777H	0.98	2.16	2.20	1.7.		1.48	3.06	1.59	3.61	
	200070	Nactar	4.49	3.27	2.20	2.38	75.	8.78	0.80	0.20	1.07	•
•	30042	207071	610	0.29	2.20	1.26	3.11		1.36	3,30	3.35	7
892	782541	AA448480		2	2.20	<u>7.</u>	2.03	B	5	1,12	1.09	8
	813641	M4774		200	2 19	2.26	3.85	3.24		0.20	5.50	7
	126568	R06862	3.87	0.40		1.87	1.73	3.63	0.20		0.98	7
695	345553	W73889	9.02	0.11	<b>6</b> (7)	-	1.74	2.81	3.42	3.40	6	8
	51947	H24313	2.05	4.48	8L.Z	2 2	0.11	3,51	0.13	1.93	<b>5</b>	. ~
289	40010	P.54038	0.09	0.21	2.19	5 5	3 14	2.20	0.53	4.87		
808	383188	AA074224		9.6	2.18	8. °	1.50	0.27	3.22	1.35	27.10	• •
	72531	T55639		0.11	2.18	3.83	20.0	6.02	3.04	0.78	58.L	• •
		A A 4 4 8 7 4 B		6.63	2.18	0.48	0.02	285	1.29	1.24	1.53	7 (
3	70007			13,38	2.18	2.11	3.23	4.	1.01	0.17	0.77	7
Ę	2000	244040	2	7.68	2.17	4.03	4.62	C#:7	2 60	0.77	2.79	7
702	46158	9/080H	, ,		2.17	3.17	3.06	60	896	3.94	1.06	7
ğ	283181			8F 4C	2.17	1.43	3.18	0.73	90.	4.03	4.23	7
ş	647985	-		2.5	2.17	1.28	0.83	127	9 :	1.18	1.39	~
705	491001	AA138710	_	14.30	. 9	1.67	1.35	3.82	G. 4.5		88.0	7
8	1481161	I AA888038		75. 75.	DL72	89	4.53	1.35	3.60	51.1	• 23	2
707	757503	AA426380	1.95	1.21	2.16		3.70	1.70	0.98	0.72	Q	
2	280375			2.51	2.10	0.4	7.	1,81	3.80	0.88	0.92	• •
2 8	283018	_		3.09	2.16	1,68	6.00	1.77	1.30	2,36	3.09	7 •
2 5	121729			1.58	2.15	45.	5 6	0.98	5.28	1.08	3.20	7
٠,	70000			2.77	2.15	1.58	B					
Ε	08080						21					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		700			1	September 123/4 Foldinph121/A Foldinph124/A Foldinph122/A Foldinph123/A Foldinph126/A # of Tumor	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	s of Tumor
	IMAGE	Accession	Avebph	AveTumor	Fold(AVe lumor/	AV FOIG[IIIF III LECT.	veBPH)	veBPH)	veBPH)	veBPH)	Vebru)	
	Cione 10	Numbor					•	133	3.02	1.85	3.18	7
712	207370	H58834	0.13	0.27	2.15	1.14	1.30	3	70.6	1.88	0.98	79
: :	307000	14306A	2.15	4.60	2.14	1.38	3.31	90.0	17.7	3	960	7
2 ;		100000	2	000	2.14	4.28	0.9 <b>4</b>	4.27	05.1		*	~
Ž.	870138	AA321354		9	2.13	3.57	3.32	2.78	0.71	05.1	? ;	
73	322481	W15305	11.7	0.4		7.7	1.13	£.,	1.68	4.18	• ·	<b>,</b> ,
718	188232	H45711	0.81	1.31	2.13		3.78	3.35	2.45	0. ¥	0.47	7
717	107525	H52001	2.60	5.54	2.13	2.40		178	3.41	1.02	0.69	84
718	480279	AA877629	0.88	<b>‡</b>	2.13	2.50	3.41	2 5	3.87	3.07	1.03	2
7	777276	N53492	330	1.12	2.13	1.10	2.23	0.5	5 6	0.59	1.88	'n
= =	82029	TRZROZ	142	3.02	2.13	0.99	3.50	3.33	7.50	3 5	76	. ~
3	700	200101		150	2.13	98.9	0.19	1.08	er.o	- 6	800	~
12	276871	N38470	3 9	;		0.40	4.34	0.31	2. 2.	90'1	3	
22	417059	W87801	0.12	0.25	3.	2 5	3 + 5	3.23	1.23	1.67	2.04	7
723	294255	N99839	13.11	27.87	2.13	1.4/		÷	1.40	4.42	<b>7</b> 6.0	8
707	243700	N45083	<del>2</del> .	2.84	2.13	1.35	3.02	9 6	 8c	1.60	2.84	7
ž	285040	NA3375	, 90, 60,	12.63	2.12	3.18	3.00	8 i	3 5	28.0	1.02	84
3		740043		2.82	2.12	1.70	3.05	14.	60.4		5.73	2
128	150	2000	3 ;	i	24	0.90	1.15	3.08	0.82	78.0		
727	743465	AA609385	A		;	3.15	3.51	2.17	1.18	1.40	7	<b>4</b>
728	154485	R54850	4.3	8.09	7.7	2 6	96.0	3.49	2.09	0.98	2.01	N
729	814780	AA455227	0.13	0.27	2.10	80.5	? ;	3.44	1.85	1.43	3.49	7
55	322005	W37418	0.91	1.80	2.10	1.29	1.14		62	1.17	1.68	~
734	418945	W87710	3.08	6.46	2.10	<b>3</b> .	3.31	B 6	2.68	3.20	1.03	7
1	R34470	AA182880	6.29	13.17	2.10	1.20	3.14	7.7	3 .	4.58	4.35	п
7	7707607	9990098		0.18	2.08	0.17	1.06	0.74	4.07		0.28	~
3	1000	Avecases		0.37	8	1.24	1.24	1.45	4.28	60.4		
ĕ	345178	W72283	9.79	6.5	3 6	70+	3.20	<b>1.6</b>	3.65	0.99	Ę.	<b>y</b> (
2	878652	AA670200	₹	3.08	60°2	<b>5</b> .	6	3.08	1.55	1.49	2.07	7
736	388511	AA026609	<b>6.7</b>	8.89	2:08	1.3/	700	3 65	0.87	1.38	1.95	8
757	235173	H73013	<del>1</del> .	2.89	2.08	1.48	3.21	9	3.12	4.93	0.84	8
738	296448	N74623	0.08	0.13	2.08	2.84	6. 10		96	1.38	4.0	7
2	357264	WR3585	1.1	2.28	2.08	2.07	3.92	28.0	900	0.28	6.10	7
	90200	042840	0.33	0.67	2.05	1.08	3.66	<b>3</b>		282	1.14	8
?	BB/B?	A 6 6 6 7 7 7 8		2.45	2,0	0.82	3.31	8.	80	4 4	9.	
Ž	/ca294	AA664423	9 :		2	4.41	2.53	<b>9</b> 1.0	0.83 83	1.10	2	. (
742	282770	N83646	0.04 0.04	0.15	5 7	,	3.58	1,85	1.80	0.32	0.80	N ·
743	195835	R89317	0.28	0.53	3	2 9	2.24	3.13	2.80	0.64	0,50	7
75	663873	AA101299	0.58	<del>1.</del>	2.03	2.13	3.2.0	86 0	323	5.47	1.68	7
\$	785522	AA454218	0.05	0.10	2.03	0.68	0.20	2				
!							22					

Table 2-1:T3NO vs. BPH; Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		•			Semple Cold(MPM121/A Fold(MPM124/A Fold(MPM124/A Fold(MPM122/A Fold(MPM125/A FOLD) VeBPH) Semple Cold(MPM121/A Fold(MPM124/A Fol	SIAIMPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A F vaBPH)	-old(MPM125/A vebph)	Sample
	u C			AveTumor	Fold(AveramonAv	(HGBPH)	veBPH)	<b>V804</b>			1.27	8
•		_	Avern		בל בל בל		•	3.98	1.13	3,33		2
		Number		•	e c	1.37	21.1		28.0	0.40	? •	•
748	42035	R58068	0.19	0.38	8 8	4.97	1.05	c.5	1.83	1.37	3.85	<b>,</b> (
	980412	N47208	0.02	0.05	3 7	5	0.21	3.77	9 9	1.83	3.68	N
: ;	10030	AA283693	<b>\$</b> 0.0	0.09	2.03	3.25	0.81	1.78	6.5	1.33	0.97	7
2	1	062630	620	0.68	2,02	9 9	3.11	1.88	3.12	2.10	3.08	7
7.78	40004		112.27	228.02	2.01	60.	3.06	1.36	1.47	9 6	7.7	64
750	1486109		20	0.55	2.01	1.00	2.5	1.08	3.17	6.5	66.0	7
751	415806	W86196	4 6	φ. •	2.01	0.73	0.0	305	3.01	<u>.</u> 33.		2
752	399421	AA733177	2.08	<u>.</u>	201	1.77	1.69		7	1.38	0.77	
753	744862	AA625907	10.80	21.69	5 6	45.5	3.53	10.8	¥.05	08:0	4.37	٠,
184	446411	W83147	0.65	<del></del>	2 6	1.48	0.12	1.12	, c	2.31	3.41	7
- 1	84404B		90.0	0.18	201	28.0	3.86	0.83	10.0	4.27	0.70	~
6	07903		0.14	0.27	2.01	2 5	0.28	1.31	9.4	1.74	0.87	7
8	2000		2.74	5.47	4.99		3.38	1.83	3.0	02.8	0.87	7
151	58805		7.33	3.4	1.99	1.20	4 06	1.89	0.82	5. J	0.48	7
758	278498		2	0.65	1.98	0.56		1.14	3.73	16.1	676	7
759	32882	R43910	3 5	28.18	1.98	1.66	5 6	1.85	3.69	1.07	2 6	•
26	878785		13.20	2.0.5	1.97	1.41	3.00	3	3.67	<u>.</u>	0.42	
781	827105	AA190834	9.88	19.43	. ¥	2.41	3.44	C ()	0.81	77.0	122	7 (
782			1.78	3.45	26.7	3.53	4.14	4.49	5	09.0	3.40	7
787	-		721	203	C8:1	1.25	3.84	1.08		1.28	1.18	7
3 7			0.13	. 0.25	ca:L	98	3.61	1.32	95.5	1.68	1.04	7
5 8			3.50	6.82	C8.L	2	3.06	0.98	80.1	0.15	15.4	2
. 6	•		0.51	0.99	1,95	; ¢	3.60	2.85	0.63	0.56	. 0.07	N
8 9			0.07	0,13	1.94	0.30	1.24	9.70	77.5	3.22	3.15	7
ē P			0.13	0.26	<b>5</b> . 1	2	0.87	1,45	3 1	3.10	3.11	~
8 8	-	_		•	1.93	0.78	1.30	1.49	07.F	3,17	3.68	7
1 2				1.13	40.1	1,31	1.88	0.78	5 6	3.99	4.78	~
					76.1	0.80	0.16	1.67	2 8	0.99	3.01	~
1		.6 AA448257			26.	4.92	0.83	0.83	9.50 7.8	0.32	0.68	~
! #					78.1	1.96	3.63	1.18	2 6	1.23	2.43	~
1					78.1	3.26	3.09	0.82	9.00 4.4	0.39	0.38	74
					1.82	090	0.39	4.58	0 6	0.92	3.71	7
<b>8</b>					787	122	0.65	1.41	5.00	0.78	1.04	~
#				•	28°1	2.14	3.20	1.12	- ¥	3.58	5.59	7
: F					98	0.18	0.16		8:-			
7	779 \$30608	08 AA071089	9 0.05	0.0			23					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession Number	AveBPH	AveTumor	Fold(AveTumoriay Fold(MPM121/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor obph) vorph) vorph) vorph) vorph) vorph) vorph) vorph) vorph) vorph)	Fold(MPM120/A veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A vebpH)	Fold(MPM122/A   veBPH)	Fold(MPM123/A vebph)	Fold(MPM126/A veBPH)	# of Tumor Samples
780	208570	H63223	0.59	1.10	1.89	0.51	0.48	2.48	3.11	3.59	1.1	7
781	131887	R25186	2.12	3.98	1.88	3.19	1.11	1.57	0.87	40.7	3.49	8
782	341978	W81361	1.28	2.41	1.88	3.07	3.06	1.81	1.31	0.70	1.31	7
783	461727	AA682293	1.49	2.80	1.88	3.10	7.00	2.34	3.24	0.89	0.92	8
28	815550	AA456818	3.83	6.87	1.87	0.83	0.89	1.54	0.50	4.16	3.30	7
285	3388	R44538	7.22	13.49	1.87	0.98	3.01	1.39	4.56	0.80	0.48	7
788	304	R44754	0.12	0.23	1.87	0.95	3.90	1.51	1.20	0.08	3.55	~
787	743189	AA401434	0.04	0.08	1.86	2.05	3.30	1.81	3.55	0.24	0.23	~
788	247194	N54036	0.36	0.66	1.88	1.09	3.23	128	3.14	1.38	1.06	7
789	183771	H4448	1.91	3.58	1.86	1.23	3.04	122	3.07	1.82	0.79	~
780	491764	AA150502	90.0	0.14	1.86	3.80	3.09	120	1,47	1.48	0.13	~
781	782594	AA447531	0.02	9.0	1.85	3.26	3.78	2.14	1.09	0.42	0.42	~
782	471859	AA035144	0.03	0.05	1.85	3.10	5.05	0.57	0.47	1.55	0.39	7
793	276712	N34895	0.13	0.23	1.85	0:30	0.39	5.27	1.75	0.10	3.28	8
767	44154	H08154	1,53	2.83	1.85	2.02	3.26	0.79	3.22	1.20	0.59	7
795	261274	H98255	2.71	5.00	1.84	1.16	3.49	1.61	3.99	0.39	0.42	~
<b>38</b>	839081	AA487608	11.79	21.75	1.84	3.03	3.73	0.71	2.01	09:0	0.98	7
797	502625	AA134576	0.88	1.61	7°.	4.66	0.80	0.64	0.68	0.19	4.08	~
798	77577	158832	0.29	0.52	1.83	1.41	3.32	1.19	3.24	1.13	0.88	~
789	839829	AA489782	0.23	0.41	1.82	1.28	3.14	1.17	1.64	3.17	0.63	7
8	53081	R15784	0.11	0.20	1.82	1.57	0.53	29:0	0.77	4.18	3.24	7
<b>80</b>	356531	W86134	2.78	5.05	1.82	3.97	0.53	1,15	0.95	0.69	3.60	7
803	32339	R42823	0.14	0.24	1.80	0.81	0.47	3.98	3.10	0.42	2.00	~
83	417688	W89074	0.20	0.35	1.80	1.48	0.57	0.05	1.03	3.31	4.34	~
춣	1435339	AA857748	9.0	0.07	1.79	0.22	3.54	67.0	1.30	0.25	4.72	~
88	128973	R10301	0.05	0.09	1.75	0.83	4.27	3.36	1.30	0.20	0.43	7
88	381323	AA017544	2.04	3.57	1.75	0.84	1.00	3.73	3.19	0.89	0.88	7
807	568440	AA148862	0.05	0.09	1.75	0.31	0.77	1.71	3,44	0.20	4.04	8
808	857198	AA773139	97.0	0.79	1.74	1.59	3.02	0.84	0.48	0.90	3.53	~
808	588858	AA130884	0.03	90:0	1.71	6.02	0.31	3.02	0.31	0.31	0.31	~
55	741891	AA401972	5.81	9.84	1.69	1.65	0.69	0.62	0.74	3.11	3.35	~
811	810027	AA455275	0.0	0.07	1.69	4.25	0.80	0.94	3.45	0.57	0.18	7
812	611609	AA455020	0.03	0.05	1.65	5.05	0.33	0.21	3.08	0.21	1.05	~
813	361607	W92431	0.21	0.35	1.63	0.45	3.01	1.77	0.79	0.33	3.44	<b>~</b>

**7** 

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

					of Tumor Fold(MPM122/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor				Fold(MPM122/A	old(MPM123/A	Fold(MPM126/A	\$ of Tumor Samples
		410				Fold(MPM120/A	Fold(MPM121)	(NABPH)	veBPH)	(appro)	!	c
	MAGE	Gen Bank	AveBPH	AveTumor		veBPH)	ALL CORN		ē	3.13	1.83	. (
_						40	0.10	0.51	90.5	0.68	3.79	7
			Ç	0.16	1.80	0.80	9	0.38	3.52	5 6	0.43	74
8 4 4	785538	AA459651	<b>&gt;</b> •	8	1.59	6.51		0.43	3. 8.	2	1.52	7
815	260168	N32071	0.05	3 6	1.59	0.28	5.10	18 6	0.45	0.16		7
4	608835	AA169154	0.02	\$ 5	15	0.21	3.55		79.0	0.18	5 6	7
	47187	H10404	900	0.10	<u> </u>	1.00	3.48		1.63	3.05	0.27	, 6
2		A A 4 59383	0.05	0.09	(6,1	3.03	0.23	1.13	0.0	0.18	<b>7</b> ,	. (
8 18 8 18	933007	A 4 100898	9.	0.0	56.	<b>88</b>	0.39	3.52		0.22	0.22	× •
<b>8</b>	00008+	900014	0.11	0.17	1.63	9 6	3.40	077	<u> </u>	0.31	0.87	Ν.
820	769579	AA423040		0.0	1,47	976	3.34	0.69	L8:0	0.33	0.33	
821	454083	AA676996		0.39	1.47	3.02	0.48	3.30	3.34	040	3.07	~
ឌ្ជ	52071	HZ4350		80.0	1,35	0.33	. 60	1,14	0.23	74.0	0.17	-
8	511066	AA100296		0.32	1.33	3.08	50.0	0.17	2.61	0.1.0	2.32	<del>.</del>
824	742818	AA400482		4 88	32.02	186.97	. 4	0.14	0.14	133.48	0.32	-
828	504184	AA152178		3 7	28.82	2.37	n (	132.15	0.22	¢. i	4.0	-
828	306420	08928N	0.04	CR.	77.47	0.60	1,38	3.48	0.75	91.20	5 6	•
A27	452374	AA700878		80.73	87.8	2.18	2.14	9 6	2.14	46.98	67.7	-
3 8	384015	AA702640	-	5.24	2 -	0.83	0.14	87.0	0.14	40.15	AC.F	• •
8 6			0.07	2	6.0	1.37	0.78	0/.L	35	0.58	33.43	- •
8			90.0		08.7	0.62	0.93	1.48	7	2.80	2.23	- ,
3				0.97	6.43	308	2.15	24.89	7.	0.25	31.39	-
<u>2</u>				1.89	6.03	06.4	0.45	0.30	es.r	0.69	0.65	<b>-</b> -
832				1.55	5.77	0.90	1.18	0.70	30.40	4 70	27.99	-
833	•	•			5.61	0.07	68.0	0.13	1.21	10	27.41	-
<b>8</b>					5.42	0.52	000	0.63	1.19	2 2	24.70	-
835	364563	-			5.40	1.15	6.6	2.14	1,43	1.59	1.74	<b>-</b>
838	3 488422	_				0.82	17.1	1.81	1.51	24.08	0.28	-
23	7 50799	H17630				2.39	8/D	28.18	0.28	80: <del>-</del>	4.0	· <b>-</b> -
88	8 302180					1.88	4.08	26.26	1.46	0.14	5 6	-
839	128083	3 R09747		_	4.96	96'0	0.82	25.48	0.07	0.08	66.0	**
28		3 R36969				2.48	0.09	30 66	1.24	1.38	0.02	-
178	•	36 H68542				0.95	1.14	6.27	2.09	22.55	0.40	. •
5 3		_		•		0.13	1.05	0.32	20.54	0.78	2.08	- •
, <b>2</b>			0.15			0.57	1.15	67° -	0.42	1.17	0.00	
. <b>ಪ</b>	844 381291	91 AA017468		6.4		19.43	2.89	23.85	0.22	0.21	- 5	. 🕶
<b>~</b>	••			·		0.47	3.0	1.32	1.01	0.69	3	
κò			479 2.41		15 4.23	19.31						
•	647 510381	81 AA055585			•		52					

**5**8

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession Number	АувВРН	AveTumor	Fold(AveTumon/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor eBPH) veBPH) veBPH) veBPH) samples	Fold(MPM120/A   veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A I	Fold(MPM123/A F veBPH)	old(MPM126/A VOBPH)	# of Tumor Samples
878	448401	AA777536	0.45	1.91	4.21	0.91	1.74	18.40	0.98	2.09	1.12	-
848	604940	AA160619	0.17	0.70	4.16	9 1,19	1.74	0.82	1.23	19.01	0.95	-
. 820	122288	T99158	0.42	1.73	4.08	1.32	2.06	1.18	1.28	2.94	15.68	-
198	781442	AA428603	0.39	1.58	4.03	1.32	1.09	0.51	0.38	20.03	0.86	-
852	430004	AA034014	023	0.92	3.97	1.74	1.54	0.57	9.0	17.98	1.99	-
853	23421	R38261	0.58	2.28	3.82	2.13	2.98	1.30	1.23	14.18	1.67	-
<b>3</b> 58	824211	AA490975	0.43	1.05	3.80	1.08	1,29	15.59	1.72	2.23	0,91	-
855	43065	R81868	0.12	0.43	3.74	1.37	0.85	2.14	0.67	14,52	2.78	-
858	347183	W80668	0.48	1.65	3.58	0.59	0.93	1.22	1.92	<del>1</del> ,22	15.62	-
857	32381	R43452	76.0	3.38	3.48	2.08	1.14	1.62	. 10.80	2.71	2.56	-
858	692778	AA158211	9.0	0.12	3.46	0.47	0.21	1.38	0.63	1.95	16.18	-
828	454808	AA877403	0.60	2.05	3.45	0.77	0.38	1.22	14.94	1.32	2.08	-
88	340641	W56760	0.27	0.91	3.41	1.52	1.27	1.69	15.08	0.51	0.39	-
861	897722	AA598983	90.0	0.26	3.40	16.70	0.81	0.76	0.85	1.03	0.13	_
862	866882	AA878352	0.19	9.0	3.36	17.69	1.01	0.60	0.60	0.21	a.ois	-
863	130664	R22579	99.0	2.22	3.35	1.17	0.97	14.33	1.21	1.19	1.22	-
<b>38</b>	1808080	AA995508	77.0	1.59	3.34	1.80	1.58	1.72	1.32	11.93	1.68	-
865	768260	AA424850	0.82	2.69	3.28	1.68	12.82	2.38	0.89	1.08	0.83	-
888	46830	H10028	0.28	0.92	3.27	1.15	66.0	0.74	1.64	13.18	1.89	-
887	1035785	AA829117	10.22	33.26	3.25	0.55	0.28	1.17	15.74	1.48	0.32	-
888	591025	AA159068	0.21	0.67	3.24	1.63	0.50	2.32	1.36	2.40	11.24	-
898	194515	R86242	0.58	1.87	324	6.47	2.02	2.78	2.94	2.78	2.43	-
2	308412	N93790	0.02	0.05	3.21	1.60	0.62	1.68	0.62	12.88	1.87	-
178	787449	A4418000	2.78	8.83	3.20	1.31	1.70	0.76	2.24	10.28	2.91	<del>-</del>
22		AA455197	0.09	0.28	3.19	0.83	0.72	0.87	0.94	0.35	15.41	_
55	322175	W37778	1.80	6.71	3.18	2.63	6.05	2.60	2.45	2.92	2.41	-
874	287770	N82251	0.10	0.31	3,14	1.79	0.93	12.98	2.84	0.10	0.23	-
875	51216	H18471	0.33	1.03	3.13	1.22	1.79	1.32	1.82	11.65	1.01	<b>-</b>
876	591683 /	<b>441214</b>	0.03	0.10	3.13	0.62	0.67	11.85	2.53	1.55	1.85	-
877	51951	H24317	0.33	1.01	3.10	1.50	12.98	1.31	1.08	0.77	1.00	-
878	_	AD04824	0.19	0.58	3.08	1.55	6.34	2.94	2.52	2.48	2.64	_
878	3448	R44164	0.46	1.41	3.05	0.95	11.69	0.77	1.46	1.05	2.37	•
8	280217	N47941	6.29	19.12	3.04	1.98	1.84	0.85	0.83	11.88	1.12	_
<b>2</b>	360638 A	AA017141	<b>5</b> .0	1.01	3.00	2.60	1.86	2.57	1.88	1.18	8.11	-

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Bank			Fold(AveTumoríav Fold(MPM121)A Fold(MPM124)A Fold(MPM122/A Fold(MPM123)A Fold(MPM126)A # of Tumor 	Fold(MPN120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A VøBPH)	# of Tumor Samples
	WAGE	•	AveBPH	AveTumor	•BPH)	veBPH)	VeBPH			3	0.81	-
		Number				000	0.96	12.90	1.25	3		-
Ç	720048	AA418543	1.1	3.32	2.98	0.80	790	13.67	0.74	12	100	- ,
3 6	705207	AA454155	2.04	6.03	2.85	0.80		168	7.41	1.89	1.85	- •
3	10000	200	0.39	1.13	2.91	2.12	7.7		1.88	7.67	1.73	-
	/67881	C//CAY		1.73	2.91	1,15	2.19	7.03	5.15	2.58	2.12	-
885	462003	AA780027	3 6		2 80	2.81	2.95	<b>3</b> .	2 5	1 88	10.26	-
888	703943	AA278145	3.01	¢/.0	9 6	1.39	1.34	1.32	1.43	3.4	2	-
887	781027	AA448031	0.83	1.82	P. 7	0.87	<u>7</u> .	0.52	2.68	A :		-
888	625516	AA187595	0.05	9.08	2.80	0.32	0.45	1,22	10.05	<u> </u>	3 7	. •
8	R18478	AA457517	4.08	11.68	2.88	1.28	7.7	2.29	127	1.75	<b>5</b> .	- ,
8 8	48387	H08859	1.41	<b>4</b> .08	2.88	2.57		2.53	1.74	7.18	1.79	- •
8 8	242763	H72322	0.28	0.73	2.87	1.57	2.40	Ė	1,73	9.16	1.31	<b>,</b>
<b>3</b>	61513	475410	0 32	0.63	2.87	0.97	2.25		138	2.07	6.39	<b>v-</b>
288	126448	2000	190	1.70	2.88	1.99	2.55	2.30	1.03	1.03	1.18	-
883	300543	COCAN	13	3.49	2.86	0.86	0.81	7771	2.15	7.24	1.48	-
ğ	25411	N22230	1 2	170	2.85	2.05	2.36	08.1	250	5.64	1.58	•
895	189285	K85883		2.40	2.84	2.50	2.20	2.20		2.87	7.85	<del></del>
888	51918		<u>.</u>		2.83	1.63	1.87	1.82	<u> </u>	2,69	2.38	-
897	111370		88.	\$ 6	2.83	1.76	2.11	0.92	7.08	8 F	2.07	-
808	376435	AA039713	1.18 8.	<b>4</b> , 5	. 88	1.10	2.13	1.85	1.85	2 7	2.	-
669	24676	R38809	0.40	1.13	19.7		0.97	10.42	1.72	1.24	14 94	. 🕶
8	127831	R08838	0.68	1.85	2.79	1.5.	670	0.62	2.18	0.57	71.6/	
Ş	77915	T61323	8.10	22.58	2.79	DO: 1	900	2.59	5.69	2.57	89. F	- ,
3 8	327481	W20438	97.0	2.12	2.78	2.05	60.0	1.45	1.03	2.33	1.61	- •
8	40283		0.15	0.41	2.78	7.84	1 6 <del>1</del>	2.83	1.66	2.95	5.61	- ,
3 2	826109		0.40	1.11	2.78	1.65	. ·	9.91	1.30	1.08	1.28	- •
5 8			9.0	8.	2.77	1.55		173	1.69	10.48	0.03	- •
3			0.35		2.77	1.16	5.7	1.80	8.37	1.28	1.29	
3 6			0.57	1.58	2.78	1.73	2:1:	0.72	1.13	0.69	4.02	
80		•	0.09	0.26	2.75	12.48	97.5 •	10.19	1,11	1,44	1.48	- •
8 8			1.43	3.91	2,74	0.94	67.	9.71	1.37	1.26	1.62	- •
3 5			0.84	2.57	2.73	1.07	<u> </u>	2.68	6.30	1.99	1.87	-
2 7			2.86	7.75	2.72	1.73	97.1	130	10.18	0.79	0.80	-
5			1.00	2.71	2.71	1.62	B 4	1.75	1.84	1.63	1.16	<b>-</b>
			0.33	0.90	2.70	8.20	3 5	2.53	6.58	1.45	0.60	-
5 4		_	•	40.69	2.70	2.31	2.12	2.02	7.45	2.53	1.28	-
915			4.02	10.78	2.68	<del>1</del> .3						
•							27					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession Number	Ачеври	AveTumor	Fold(AveTumoriav, Fold(MPM12DIA Fold(MPM121/A Fold(MPM122IA Fold(MPM123IA Fold(MPM125IA \$ of Tumor ebpt) veBpt) veBpt) veBpt) veBpt) samples	Fold(MPM120/A   veBPH)	Fold(MPM121/A	Fold(MPN124/A veBPH)	Fold(MPM122/A veBPH)	Fold(MPM123/A I	Fold(MPM125JA veBPH)	Samples
916	44303	H06380	1.89	5.32	2.68	1.57	2.81	<u>7</u> .	1.80	5.50	2.76	-
917		W01211	0.81	1.63	2.68	1.80	6.53	2.29	1.39	1.88	2.10	-
918		R64231	0.12	0.31	2.67	1.98	7.39	1.13	2.15	1,48	1.91	-
919	608228	AA179161	0.09	0.25	2.67	0.51	2.46	0.93	1.38	0.33	10.39	-
920	•	AA609368	0.58	1.64	2.84	1,47	1.50	8.81	1.78	1.27	1.02	-
921		N62948	0.33	0.87	2.63	2.06	6.78	1.41	2.87	1.46	1.20	-
822	505338	AA147594	0.08	0.25	2.63	2.25	0.11	2.32	0.83	1.71	8.55	-
923	63022	R15880	0.89	2.34	2.62	2.58	1.25	1.32	6.01	1.69	2.88	-
824	273546	N33274	2.51	6.55	2.81	2.25	2.58	2.24	8.64	0.98	1.01	<b>-</b>
928	843048	AA485983	0.10	0.25	2.61	2.59	1.1	8.50	127	0.72	1,45	-
858	594517	AA169645	31.25	81.32	2.80	1.36	2.19	2.32	6.51	1.76	1.48	-
128	453175	AA700218	0.99	2.58	2.60	5.15	2.24	1.63	2.78	1.71	2.03	-
828	380357	AA013355	0.26	0.87	2.80	1.94	9.92	1.20	1.48	0.37	0.70	-
929	345525	W72437	3.95	10,21	2.58	2.40	2.95	1.23	5.65	1,74	1.53	-
83	344274	W73811	0.13	0.33	2.58	0.08	69.0	12.77	0.63	0.28	1,15	-
931	389152	AA733105	0.03	90.0	2.58	8.38	0.82	1.72	2.27	0.98	0.31	-
832	787345	AA418584	1.03	2.65	2.58	1.48	1.68	2.03	2.47	6.56	2.28	-
88	743727	AA628355	98.0	272	2.58	0.98	2.48	5.63	1.51	1.91	2.98	-
8		AA484755	0.07	0,18	2.57	1.64	1.73	1.89	1.42	7.19	1.67	-
836		AA872034	0.20	0.51	2.57	1.98	1.32	1.27	7.42	2.20	1.24	<b>-</b> -
838		AA498252	0.87	2.23	2.57	1.20	1.20	1.67	8.80	0.59	1.98	-
937		R45157	0.59	3.	2.57	1.13	1.11	1.07	0.94	0.72	10.42	-
808	_	AA701519	0.50	1.29	2.58	2.29	1.57	2.64	1.01	0.64	7.34	-
838		R22003	3.70	9.47	2.58	1.96	5.20	1.35	2.98	1.57	2.33	-
3	30428	R42061	3.15	8.00	2.54	1.75	2.23	0.74	2.28	2.59	5,66	-
2	299162	N75473	0.03	0.07	2.54	0.38	0.38	2.96	9.96	0.38	1.25	-
85	267879	N25552	0.58	1.47	2.53	2.42	5.39	1.41	2.09	1.48	2.38	<b>-</b>
Z	145383	R77919	0.10	0.24	2.52	1.78	1.91	1.10	1.86	8.26	0.17	-
3	154720	R55220	0.05	0.13	2.52	2.29	7.29	1.91	0.19	1.86	1.60	<b>-</b> -
3	395853	AA757564	0.49	1.22	2.52	1.20	1.50	1.33	0.70	9.11	12	-
Z	412949	AA707840	0.92	2.31	2.51	1.63	6.07	2.31	1.74	1.69	263	-
ž	120892	T95898	3.38	8.48	2.51	1.12	1.30	2.13	1.84	1.88	6.82	-
Z	271989	N31935	0.18	0.45	2.50	1.45	2.91	1.52	1.12	5.41	2.57	-
ž	592592	AA158577	3.38	8.37	2.49	1.59	0.54	2.12	9.45	0.77	0.49	-
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Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

		Gen Benk			polytavotumorjav foldinpnizoja foldinpnizija foldinpnizaja foldinpnizzja foldinpnizzja foldinpnizsja 8 of Tumor	Fold(MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A	Fold(MPM126/A	# of Tumor
	Clone 1D	Accession	AveBPH	AveTumor	•BPH)	vaBPH)	veBPH)	veBPH)	(HGB4)	VeBPH)	VeBPH)	Seidules
Ş	E04068	4444407	800	0.20	2.49	7.00	1.24	2.42	1.57	1.07	1.62	-
3	ecollec	ANIOIAN MAIOIAN		2 6	2.48	6.02	2.69	1.38	1.52	1.97	1.31	-
	707/07	N445/8	3 -	27.5	2.48	1,34	1.22	0.91	1.08	1.21	9.12	•
	337048	C+7017	. 4	65.0	2.47	9.38	1.43	0.13	1.53	1.88	0.48	
3 3	304K44	NROADK	0.51	125	2.47	2.70	5.07	2.03	2.61	2.0	<del>.</del>	<b>-</b> ·
S &	202004	H44020	28.65	70.78	2,47	2.13	1.92	1.40	5.51	1.96	1.82	<b>,</b> - ·
3 3	119458	T00074	0.27	0.67	2.47	1.93	2.00	1.57	1.15	5.60	254	<b>,</b>
3 5		A 4 8 1 7 K 4	25	0.81	2.47	1.81	1.94	1.23	1.48	6.78	1.58	<del>-</del> ·
8 8	411033	AARTRARA	1 1	2.73	248	1.61	5.16	2.27	1.48	2.27	1.98	-
	350853	401010101		980	2.48	0.87	2.24	2.13	1.48	1.27	6.61	-
	200000	AA358487	9 6	3 5	2.48	5.	2.61	86.	6.29	1.80	2.03	<del>-</del>
8 8	340400	67000	5 6	2.25	2.48	0.90	0.76	10.09	1.06	0.89	1.07	•
5 8	60100	aren a	5	0.78	2.46	2.80	7.68	0.56	0.87	1.05	1.89	
8 8	3 2 2	27276	, E	88	2.45	2.77	1.25	0.82	1.18	2.92	5.77	<b>-</b> -
2 3	846773	AA484751	, e	2.5	24	1,12	1.19	8.67	1.17	1.26	<u>1</u>	<b>-</b>
ğ	2000	N54407	50	1.29	2.44	2.00	1.80	5.73	1.70	1.87	1,43	-
3 8	201304	27.000	2.54	6.16	2.42	1.35	2.68	2.98	5.20	1.37	98:0	<del>-</del>
} }	773470	AA428514	0.17	9	2,42	1.55	1.08	1,04	0.90	8.27	0.68	<b>-</b> ·
8	ROZRAR	AA406708	5.28	12.74	2.42	2.58	6.50	0.47	1.05	1.75	2,16	<del>-</del> ,
8	111884	TAKRAR	0.03	0.07	2.41	1.89	1.77	2.00	1.45	2.36	5.02	<b>-</b>
22	744897	AAR25784	169	80.	2,40	1,57	1.49	2.43	5.19	2.25	1.48	<b>,.</b>
7.	48411	H09164	330	7.91	2,40	1.72	1.78	2.87	5.14	1.19	4.88	-
	704784	A 4 4 6 7 7 9 0	8	1.42	2.40	1.08	1.51	5.80	2.12	2.35	1.52	-
67.0	1031273	AABOBOSA	0.38	0.81	2.38	1.15	7.94	1.69	0.99	0.00	1.70	<b>.</b>
720	46129	H08529	0.07	0.18	239	0.81	1,22	1.86	0.82	6.90	2.65	<b>-</b> ,
7,0	220389	HRRAIR		2.65	2.39	1.00	1.09	1.18	1.48	7.40	2.27	-
878	1292170	AA705819	\$	3.50	2.39	1.36	1.60	8.89	1.05	0.54	9.88	<b>.</b>
6	358711	W84580	0.78	88.	2.38	1.32	2.38	1.02	1.38	6.51	1.88	-
82	4308A	A A R 7 R 2 M 3	0.65	1.56	2.38	1.54	2.21	2.38	5.04	1.26	<b>78</b> .	<b>-</b>
2 6	35575	R45964	2.28	5.43	2.38	1.71	0.81	2.32	2.25	1.78	5.28	<del>-</del> -
8	67078	40,71	0.53	1.28	2,37	0.80	1.53	1.02	8.43	1.41	2.	-
3	128205	AA844141	0.30	0.72	2.37	0.89	0.90	0.69	0.41	10.29	<del>.</del> 8	-
ŝ	197703	R93729	0.76	181	2.37	1.39	1.02	8.38	2.45	0.70	87	<del>-</del> .
<b>8</b>	26736	<b>K38691</b>	7.65	18.11	2.37	0.64	1.43	1.12	99.9	2.63	£.	-
							29					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

INAGE Clone ID Clone ID 1 39888 276815 121615 828303 3442	Gen Bank Accession Number	Ачеврн	AveTumor	Fold(AveTumorfav Fold(MPM120/A Fold(MPM124/A Fold(MPM122/A Fold(MPM123/A Fold(MPM123/A # of Tumor ABPH) varBHI varBHI varBHI varBHI varBHI	Fold(MPM120/A F	old(MPM121/A F	old (MPM124/A F	Fold(MPM122/A F veBPH)	od(MPM123/A Fo	old(MPM126/A veBPH)	# of Tumor
1 39899 1 276815 1 430007 1 121615 826303 3442				*				***			
1 278815 430007 121615 826303 34442	R52824	090	1.41	2.36	1.26	96.0	1.27	8.89	1.09	0.62	-
121615 121615 828303 3442	N34824	0.13	0.32	2.36	7.54	0.57	2.63	1.43	1.70	0.30	-
121615 828303 34442	AA034115	0.48	1.14	2.35	6.87	2.92	0.89	0.84	0.87	1.75	•
3442	T97592	77.0	<del>2</del> .	2.35	7.31	1.59	1.65	1.23	1.13	1.21	-
3442	4A521018	0.54	1.28	2.35	1.29	1.46	1.13	1.43	7.48	1.30	-
	R44985	1.70	<b>6.</b> 9	2.35	0.88	0.88	2.43	8.64	0.85	0.41	-
742542	AA400010	0.20	0.48	2.34	1.7	2.12	1.52	1.14	5.44	2.13	<del>-</del>
327078	W23758	0.30	0.70	2.34	1.23	1.45	1.58	5.33	1.78	2.68	-
160656	H24854	0.07	0.17	2.34	0.52	0.74	76.0	1.18	122	9.40	-
265114	N20820	0.3 <del>4</del>	0.80	234	7.86	0.89	1.69	1.01	0,44	2.14	-
926842	AA181424	1.69	3.95	234	1,72	1.97	2.52	0.59	0.96	6.26	-
291403	N87782	9.0	0.10	2.33	5.38	27.0	223	1,4	2.12	2.12	-
240318	H89795	0.43	8.	2.33	1.83	2.67	1.04	5.64	1.01	1.19	-
252684	H88334	<u>1.6</u>	3.82	2.33	0.77	0.58	9.21	0.83	68.0	1.48	-
302190	W16724	0.18	0.41	2.33	1.08	1,73	2.18	1.05	8.70	1.23	_
130742		0.83	<b>1.</b>	2.33	1.55	1.60	7.25	1.31	1.29	96.0	-
884883	AA629911	1,31	3.05	2.33	0.67	0.64	0.61	9.75	1.41	78.0	-
427883		0.28	0.65	2.32	1.41	1.54	1.91	1.41	1.54	6.13	-
281241		0.85	1.97	2.32	2.51	5.63	1.48	1.28	1.99	20.1	-
431028		90.0	0.13	2.32	5.12	1.92	2.46	2.71	0.8ġ	0.86	-
381996		0.46	1.07	2.32	1.41	0.59	0.79	9.48	0.33	1.34	-
629498	_	2.16	4.99	2.32	0.93	1.17	6.26	1.19	1.30	<u>5</u> .	_
3244		0.03	90.0	2.30	1.84	0.79	1.98	2.04	2.01	5.13	<b>-</b>
428778	AA005228	0.25	0.58	2.29	0.96	0.54	0.17	2.06	9.31	0.70	-
1030738	~	0.54	1,23	2.29	1.63	1.43	7.27	1.22	1.36	0.83	-
113600	_	18.96	43.36	229	0.95	1,38	1.79	9.6	2.82	0.65	-
305809		0.92	2.10	2.28	0.83	1.72	1.21	6.27	2.14	2.50	-
30502		0.11	0.28	2.28	. 5.01	2.64	2.00	1.62	1.22	1,18	-
897791	_	99.0	1.55	2.27	0.80	0.89	1.89	1.85	6.37	1.85	-
430022	Φ.	0.40	0.92	777	0.70	1.27	1.75	2.28	5.09	2.67	-
46238			0.38	2.27	5.57	<del>1</del> .	2.71	1.32	0.42	2.18	-
703739 /			19.24	2.27	0.85	0.71	0.71	1.69	0.82	9.05	-
342720	_		0.98	2.27	0.42	0.02	0.87	1.58	9.54	1.19	_
1017 25156 Ra	R38935 1	51.1	2.81	2.27	1.08	5.81	2.33	2.50	0.73	1.15	_
					8						,

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

					Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM121/A VeBPH) veBPH) veBPH) veBPH) Samples	old(MPM120/A	Fold(MPM121/A	VeBPH)	VeBPH)	veBPH)	veBPH)	e and units
- 1	IMAGE		AveBPH	Ave Tumor	•BPH)	VeBPH)			,	770	6.54	÷
_		Number				18	2.38	0.81	1.73	; c	2.58	<u>-</u>
-	754654	AA411807	0.08	0.18	2.20	506	0.54	1.89	0.13	, . , .	0.17	-
	377671	AA055979	0.08	0.17	2.25	3 6	0.35	0.38	11.15	71.1	47.	**
		A134BOR	2.29	5.16	2.25	0.33	1 65	0.93	6.21	241	5.5	• •
	RA/L/Z	20105		11.36	2.25	0.83	6.		5.99	1.42	0.89	
	448885	AA777870			224	1.38	0.94	2.63	1.28	28.8	4.0	Ψ-
<u> 1</u>	171838	H19068	0.37	\$ 1		0.48	1.43	3.	07.1	29 1	1.65	-
1023	378709	AA778089	121	2.70	<b>3</b> 1	44.0	1.29	8.10	2.62	400	1.20	-
7605	325012	AA284283	0.09	0.21	2.2	1 0	0.92	1.02	1.27	C7:1		-
, ,	10.70F.	AA019316	0.22	0.48	22	7.57	14.	1.00	1.35	1.81	10.6	-
3 5	4004	H10208	97.0	1.73	2.23	6.51	. 4 	1,02	6.04	1.07	7.37	
	1000	B7074	0.38	0.85	221	0.63		5.18	1.73	1.38	17.1	- •
30 5	13067	AA47R257	90.0	1.18	2.21	7.60		0.70	1.05	10.80	0.55	. •
1026	DAT!	1000	800	0.20	2.21	0.08	5.0	84.0	1.80	<b>1</b> .64	2.16	- ,
1929	71730	101228		200	2.20	1.59	5.46	3 5	9.15	0.27	0.27	<b>-</b>
휺	281571	H98756	3 1	, F	2.20	99.0	0.35	70.7	7.0	7.63	1.13	<b>-</b>
1031	814053	AA465495	10.76	7.37	2.20	1.24	1.46	<b>1</b> .02	- 6	0.88	1.59	-
1032	1292887	AA776755	0.48	co.T		1.76	<b>1.8</b>	5.61	00.1	4.47	130	-
£85	324660	AA284278	0.25	0.5 75.	7.50	0.86	1.78	1.92	5.87		521	-
£ 5	72000	152320	4.01	8.8 1	2.20	1 91	2.16	1.40	0.97	8.	85.0	-
\$2	1467109	AA883170	0.57	1.28	2.19		1.13	1.73	0.09	<b>5</b>	3	-
5	244051		0.25	0.55	2.19	0.0	990	8.52	0.83	0.78	700	
2 6	40.70RO	AA165512	1.86	4.07	2.19	7.0		1.89	1.69	<u>¥</u>	24.0	-
3 5	268040		1,50	3.29	2.19	3.5	120	0.81	0.50	0.22	10.5/	• •
3 5	481871		0.18	0.35	2.18	60.0	7.28	6.35	1,41	1.27	10.1	. <b>.</b> -
3 3	705588	AA459692	0.79	1.70	2.18	C.,	0.53	2.15	6.01	1.14	1.70	•
3	166199		99.	0.11	2.18	0C.1	74.0	8.54	0.12	0.12	7.7	•
3	46183		0.08	0.18	2.18	0.0	1.47	1.24	1.70	4.05	6.5	. 🕶
3	344806		0.42	0.91	2.18	1.20	0.79	0.98	1.07	0,55	5 6	
3	K4030		0.42	0.91	2.18	/A'O	6	0.15	0.19	0.88	• •	•
\$ \$	-		0.05	0.11	217	10.72	55.5	3	0.82	6.45	2 :	- <b>-</b>
3	•	_	0.51	1.09	2.17	1.57	100	0.25	0.25	0.25	777	- •
3	42027			0.09	2.17	2.12	1.51	1.37	1.17	5.78	2.04	- •
3 3				1.52	2.18	1.47	1.12	0.74	0.85	5.42	2.24	- •
3			1.73	3.73	2.18	2.42	07:-	1.55	0.48	0.57	15.8	- •
2 5			0.23	0.49	2.16	0.42	0.58	1.24	0.68	0.70	1.32	-
			28.0	1.37	2.15	1.84		•				
5							31					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	IMAGE Clone ID	Gen Bank Accession Number	Ачеврн	AveTumor	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A \$ of Tumor •BPH) veBPH) veBPH) Semples	Fold(MPM120/A F veBPH)	Fold(MPM121/A   veBPH)	Fold(MPM124/A veBPH)	Fold(MPM122/A   veBPH)	Fold(MPM123/A   veBPH)	Fold(MPM126/A veBPH)	Samples
1052	131626	R23735	0.10	0.22	2.15	0.63	1.32	1.95	1.79	6.30	0.92	-
1053	32663	R43319	0.50	1.07	2.15	0.89	1.42	1.26	7.30	1.08	0.88	<b>-</b>
105	1031717	AA609584	1.03	222	2.15	1.83	1.79	1.80	5.23	1.14	1.10	<b>-</b>
1055	254328	N22272	. 1.39	2.97	2.15	0.66	0.43	9.65	0.73	. 62.0	0.60	-
1056	788805	AA452897	99.0	1.42	2.14	1.48	2.24	2.10	5.53	0.52	<b>1.0</b>	-
1057	186245	R88267	0.03	0.07	2.14	0.31	5.65	0.31	2.51	1.42	2.64	P.
1058	32991	R44770	0.28	0.65	2.14	10.49	0.04	1.10	0.61	0.20	0.38	<b>y-</b>
1059	712840	AA282023	0.13	0.28	2.13	1.31	0.35	<u>.</u> .	0.49	8.11	1.08	-
1060	359153	AA010138	0.47	8.	2.13	1.35	1.22	6,15	1.71	1.37	0.82	-
1061	207884	H60581	0.42	0.89	2.13	1.08	1.48	2.05	1.23	1.34	5.57	-
1062	395711	AA757784	0.10	0.22	2.12	1.66	5.61	1.05	1.68	1.38	1.40	-
1063	78946	T61782	3.51	7.43	2.12	1.23	0.53	1.70	1.68	1.15	6.41	•
<u>\$</u>	42373	R87147	2.61	6.62	2.11	2.11	0.91	1.90	1.25	5.7	5.19	<b>-</b>
1065	199828	R96598	0.18	0.37	2.10	1.07	1.40	0.88	1.69	0.70	6.89	-
1066	29828	R42331	1.30	2.72	2.10	2.03	0.72	0.48	2.13	1. 1.	5.32	-
1067	809828	AA455521	99.0	1.39	2.10	6.79	1,73	0.53	1.28	0.99	1,30	-
1068	280561	N62378	2.80	5.88	2.10	1.31	1.57	0.58	1.09	0.95	7.10	-
1069	781088	AA430002	0.62	1.31	2.10	1.23	0.99	8.61	1.58	1.28	0.82	· <b>—</b>
1070	282384	N51987	0.51	1.08	2.09	1.05	0.99	1.80	1.23	1.17	621	-
1071	782873	AA447585	0.21	17.0	2.08	0.47	1.67	0.47	1.20	7.98	69.0	-
1072	73785	T64872	1.35	2.79	2.07	1.21	1.21	1.49	5.44	1.88	122	-
1073	730124	AA412470.	0.22	0.44	2.05	2.83	1.29	0.05	2.06	5.24	0.73	-
1074	51483	H18924	0.28	0.57	2.05	1.36	0.99	0.32	1.09	7.73	0.79	-
1075	38487	R48124	0.34	0.70	2.05	9.0	1.05	0.28	0.36	9.67	650	· June
1076	1048723	AA620642	0.71	<del>1</del> .	2.03	1.67	1.45	1.18	1.05	6.22	0.63	-
107	141815	R70685	8.27	18.77	2.03	0.45	0.67	0.71	0.58	8.88	0.91	-
1078	757337	AA437099	1.28	2.60	2.03	0.60	0.98	0.58	2.80	1.88	5.54	
1079	255659	N27841	0.05	0.09	2.02	7.29	1.20	0.85	0.79	1.68	0.22	-
1080	_	AA001878	0.20	0.41	2.02	0.05	0.31	0.75	0.85	9.69	69'0	-
1081	•	AA599102	2.48	5.00	2.02	1.00	1.18	6.48	1.57	0.85	0.92	-
1082	•	AA481208	0.09	0.18	2.01	0.11	1.13	0.82	1.60	8.82	1.39	-
1083	_	AA864875	12.97	28.09	2.01	1.29	1.62	1.39	5.81	1,46	0.49	<b>-</b>
5 2	•	AA443838	0.49	0.98	2.01	0.95	1.21	0.63	0.37	8.19	0.71	-
1085	252830	H88261	1.63	3.26	2.00	1.12	0.93	7.24	0.89	. 29.0	96.0	-
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ğ .						•																															
f of Tun		•-	-	-	-	-	•	-	•	-	-	-	•	-	•	•		- +-	• •	•		•		_	-	-	-	-	-	_	-	-	-	-	-	-	
cold(MPM125/A	Very v	1.50	1.27	0.99	96.0	83.0	200	6.35	6.37	1.47	80 0	0.49		4 O	4 6	0.00	6.63	3,5	6.5	2.5	0.08	6.5	0.85	0.48	2.40	0.83	0.33	0.85	0.72	0.60	1.79	0.25	1.29	1.02	80.0	0.73	}
old (MPM123/A	veBPH)	0.59	1.33			\$ 6	1.30	0.57	030	0.52	700	* 1. / 1. /	7.29	9.0	1.50	1.63	0.57	8.19	2.74	1.27	1.48	0.32	0.13	21.1 84.6	0.10	800	, t	9 08 0	5.50 \$1.50	7	1.57	7	, 6 - 6		0.1	9000	<u> </u>
Fold(MPM122/A F	veBPH)	0.85	9 9	0.00	£.18	1.57	6.15	1 78		/6'1	5.10	0.83	1.65	5.73	1.18	0.89	0.57	0.39	2.48	5.88	5.67	0.51	1.28	0.87	5.68	0.98	90.00 00.000		1.19	8 1	0.52	0.0	4.99	0.62	1.43	1.31	5.
EcidiMPM124/A F	veBPH)	5	32.5	<del>2</del> .	6.39	12:0	1.86	3 8		1.28	1.67	0.27	0.97	0.53	5.48	1.10	15.0	0.75	0.81	2.31	1,48	0.53	1.57	7.48	90.0	5.04	0.90	1.22	0.60	1.28	1.07	0.49	1.45	0.98	1.34	2.60	5.72
479000000000000000000000000000000000000	Fold(MPM121/A vebPH)		1.50	1.00	1.05	160		5.	0.48	0.87	1.70	2.83	0.61	1.40	1.30	88 0	9 c	0.65 85	30	0.73	1.38	0.68	0.75	0.58	2.45	0.89	2.20	0.03	1.75	1.29	5.94	0.76	5.80	6.61	1.11	1.33	0.90
	old(MPM120/A 1	(m. Lday	5.45	1.13	4 +	2 6	 09:-	1.05	2.00	1.33	1.5	13:1	6.30	6.7		95.T	1.40	2.53	0.63	21.0	0.00 80.00	0.00 0.000	9.97	65.0	2 63	1.12	1.50	0.89	9.9	1.23	2.65	0.82	0.25	0.85	5.24 5.24	0.27	
ESIGNEMIZZIA FOLDINEMIZZIA FOLDINEMIZZIA FOLDINEMIZZIA FOLDINEMIZZIA FOLDINEMIZZIA FOLDINEMIZZIA FOLDINEMIZZIA	Fold(AveTumor/Av	•BPH)	60		207	2.00	1,99	1,99	ee e	200	08.1	1.96	1.95	1.93	1,95	1.95	1,94	1.94	<b>4</b> .	1.93	1.93	1.93	1.92	1.92	Ā .	5 C	Q8.	1 89		2	9 2	9 6	8 8	3 5	78.	1.8/	8
	Accentioner	AVE DINO		4.16	50.73	<del>7.</del>	4.66	74 47		15.78	90.0	1.12	0.23	0.69	0.40	66.0	1.66	0.03	0.40	0.14	20.23	6.25	0.34	0.18	23	0.22	0.97	0.19	0.48	76.0	9 G	P	0.87	0.08 0.08	2.26	0.52	0.35
	i	AveBPH	F	2,08	25.34	0.73	234		9.00	7.99	0.03	0.57	0.12	. 0.35	0.21	0.51	0.85	0.02	0.21	0.07	10.48	325	0.18	60.0	121	0.12	0.51	0.10	0.26	4.72	0.38 0.38	5	0.38	<b>5</b>	5	0.28	0.19
		ç	Number	R39069	A1005519	R00809	A A B B E 3 4 4	Assessi	AAB84713	T62179	R61374	AA778839	AA428198	AA136899	R56604	N29638	A A 4 98887	TTT729	N34945	R80235	AA425438	H80325	AA491227	AA480353	AA456968	H08397	W85050	R\$2085	R44530	<b>A634945</b>	AA406228	W73894	AA400329	AA679180	H02550	H44032	N33865
	in Carrie			24918	1637296	122570	, ,		1487207	79022	37885	452545			41074	287342	807505	108378	276911	147075	773331	241077	824074	796078	838616	45501	415279	40299	33894	471641	742887	346523	743228	868702	150798	183558	243905
	=	ے ت ا	)	980	1087			1689 -	1080	1091	_											1 2	<u> </u>	105	100	1107	1108	100	1110	1111	1112	1113	114	115	1118	1117	118

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

٠.	IMAGE	Gen Bank Accession	AveBPH	AveTumor	Fold(AveTumod'Av Fold(MPM120/A Fold(MPM121/A Fold(MPM122/A Fold(MPM123/A Fold(MPM126/A # of Tumor	old(MPM120/A F	old(MPM121/A F	old(MPM124/A F	old(MPM122/A F	old(MPM123/A Forest)	old(MPM126/A	# of Tumor Samples
		Number			(uuga							•
1120	243321	H94720	1.08	2.00	1.88	0.82	0.98	1.02	5.71	<b>3</b> .	60.0	-
1121	243803	N49725	90	1,19	1.88	1.16	1.11	1:1	1.15	.05 5	6.57	<del>-</del>
1122	841610	AA487457	90.0	0.11	1.86	1.70	5.30	2.13	0.17	1.87	0.17	<b>-</b> -
1123	80371	785785	177	2.24	1.85	0.50	6.17	<u>₹</u>	0.72	1.21	1.09	•-
1124	731029	AA421289	0.08	0.15	1.85	1.68	2.25	0,69	6.70	0.12	0.67	-
1125	284882	N86737	0.33	0.61	1.85	0.44	0.82	99.0	5.74	2.61	0.92	<b>-</b>
1128	1031009	AA635188	28.2	10.78	1.85	0.60	0.46	1,52	7,50	0.73	0.28	-
1127	1030837	AARD8770	0.76	¥.	1.85	1.20	1.32	22	1.03	1.16	1.13	-
1128	273168	N36794	0.15	0.27	181	1.65	0.62	9.0	2.12	5.94	0.07	❤.
1129	66317	T68818	8.85	16.25	18.	6.02	2.40	0.49	270	0.60	<del>1</del> 28	-
4130	253931	N22084	÷	2.03	1.83	0.86	9.0	6.40	1.04	0.80	0.92	
131	487088	AA045300	2.58	4.69	1.82	0.88	0.91	5.53	1,44	1.09	1.05	-
1132	253598	H89378	1.38	2.50	1.81	0.86	0.87	5.81	0.82	0.75	89.	-
1133	1058188	A4821001	0.88	25	1.79	0.96	1,47	5.27	1.28	1.20	0.61	-
7	1606837	AA898028	31.95	57.14	1.78	1.13	0.53	1.83	5.43	0.98	0.62	-
1135	343731	W89213	0.0	0.11	1.78	0.21	5.41	1.10	2.08	0.17	1.7	-
1138	25058	R38894	60.0	0.15	1.78	0.12	0.55	1.10	7.92	0.89	0.12	-
1137	257387	N30704	2.69	4.78	1.78	0.89	1.40	1.21	5.01	1.13	<b>5</b> .	-
1138	128058	R09729	1.13	2.00	1.77	0.80	0.84	6.31	0.75	1.01	1.13	<b>-</b>
1139	397604	AA701034	0.18	0.32	1.78	0.81	0.97	0.61	0.83	6.53	0.83	
140	42271	R60328	0.02	9.0	1.76	2.08	0.50	0.88	0,50	0:20	6.13	
=======================================	71674	T57920	0.24	0.42	1.78	0.04	7.34	1.28	0.30	0:30	<del>4</del>	-
1142	340949	W57818	0.17	0.30	1.74	0.48	0.91	5.79	2.19	1.04	90.0	-
1143	788258	AA45409B	0.10	0.17	1.73	0.19	5.20	1.99	0.10	2.82	0.10	<b>-</b>
14	142397	R69834	0.66	1,13	1.72	0.90	1.00	1.08	0.97	0.50	6.93	-
1145	66902	T69477	0.89	1.63	1.72	0.49	0.39	0.57	6.34	1.28	8	<b>-</b>
1146	28508	R20640	0.47	0.80	1.72	0.96	0.75	1.23	5.01	1.66	17.0	-
1147	489476	AA045673	0.11	0.20	1.72	0.86	60.0	0.31	0.90	6.83	ឆ	-
1148	785446	AA453616	0.12	0.20	1.72	1.17	2.30	5.68	0.97	0.09	60.0	-
1149	471408	AA035347	90.0	0.14	1.71	0.44	5.02	1.21	2.24	0.85	0.51	<del>-</del>
1150	745001	AA626011	1.43	2.46	1.71	0.67	1.30	1.30	5.72	0.93	0.35	-
1151	609155	AA178857	2.10	3.59	1.71	0.49	1.4	0.60	98.0	6.09	0.78	-
1152	796549	AA480274	0.01	0.02	1.70	0.72	0.54	5.00	2.50	0.72	27.0	-
153	363003	AA018866	0.40	0.68	1.70	1.05	1.03	09.0	0.60	5.68	<u>t</u>	-
						34	_					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	MAGE			Anofinance	Fold(AveTumor/Av	Fold(AveTumor/Av Fold(MPM120/A Fold(MPM124/A Fold(MPM124/A Fold(MPM124/A Fold(MPM120/A	A FOIGINITION LAW	veBPH)	veBPH)	veBPH)	asimbies asimbies
•		=	AVeligh	Average	<b>BPH</b>			,	. 4	0.14	-
-		Numper		-		478 0.14	1.28	0.83	0.0	91.0	<b>4</b>
131	271483	N31808	0.0	0.12	1.69		1.14	5.86	0.92	D !	. •
	48758	H09938	2.53	4.28	1.69		2	1.35	0.70	٠. م	<u>.</u> .
3 5	2000	A A E O 4 3 K 4	1.09	1.83	1.68		8 6	0.41	0.91	5.18	-
	104079	10000	20	55.0	1.69	1.71	5 6	280	0.78	0.50	-
	1291950	AA707450	3 6	9	1.86	0.68	6.32	3 5	790	0.62	-
1158	418197	W80384	96.0	8	9	1.52 0.76	1.20	ar.o		0.18	-
1159	133358	R26859	0.61	<u>5</u>	3 5	0.27 2.45	0.38	7.	9.50		-
1180	269606	N26769	0.08	0.10	CQ'L		1.02	0.23	0.68	0.0	. •
	4293448	_	0.46	0.78	1.65		0.39	1.09	7.12	0.50	- ,
	700053		2.45	4.03	1.65		17.0	0.23	0.16	7.41	-
70 :	300	07027	80	0.10	1.64	0,16	9 9	0.74	5.35	0.35	<del>-</del>
=======================================	9187/	AMA/8814		0.37	1.82	2.56 0.27	o :	76.0	0.97	9.3 7.0	-
184	31869	R43017	7.0		1.62	2.59 5.16	<b>7</b> .0	5 6	2.20	0.97	-
1185	84141	171042	0.03	60.0		0.64 0.38	0.53	LO:G		5	-
1168	969636	AA683582	1,87	3.03	70"1	2000	1.15	7.87	ê .		•
1187	810897	AA457696	0.14	0.22	1.01		0.60	5.00	0.14	0.68	
	277.74	R35283	0.07	0.11	1.61		0.24	5.63	0.11	2.13	<b>,</b>
901	5666	CONCE	0.24	0.38	1,59	1.04	7	0.82	5.54	0.10	•-
1168	201001	000000		0 18	1.59	1,13 0.87	67.1	90 0	5.69	0.09	<b>-</b>
1170	112858	191168	2 ;	9	8	0.29	0.70	9.0		5.73	-
1171	786117	AA460961	0.11	2 :	3 5	133	0.85	0.11	- :	7	-
1172	41321	R56906	0.0	0. 7.	/c'1		0.32	1.31	0.75	r 6	•
1173	429448	AA007619	000	0.48	1.57		1.61	0.09	0.09	00.0	٠ •
4474	882571		0.11	0.17	55.		0.73	0.61	0.40	5,41	- ,
44.44	448813		1.22	1.89	1.55		0.78	6.22	0.22	0.33	-
	89607	•	0.22	0.34	1.54		0.48	0.88	6.51	0.31	-
2 ;	1000	A A 4.8.0.8.10	0.38	0.55	1,52			0.73	0.70	0.66	•-
	60100		191	2.89	1.51		9 6	0.31	0.31	0.31	-
8/11	20000		0.03	0.05	1.51	0.31 7.25		0.88	0.62	0.27	•
1178	0.7300		9	0.14	1.50	6.53 0.59	- :	1 6	5.78	0.55	-
2 2 2 3	128240		6 6	0.14	1.50	1,61 0.07	0.4		0.65	0.71	-
1181	22885			, ,	1.48	0.77	5.33	5	. 18	0.31	-
1182	288404		તું : -	777	7.	90.0	. 0.73	<b>X</b> .		86.0	•
1183	366915	AA027147	9.18	0.23	<u> </u>	9040	1.35	0.20	0.20	640	. •
1184	591485	AA160695	<b>0</b> .0	9 9 9	7.42		0.24	0.82	7.0	0.63	- ,
1185	344585	W73834	0.37	0.52	<del>7</del>		. 0.31	0.31	9.8 78.	6.29	- ·
1188		AA430042	003	9. 50.	1.38		0.98	0.73	0.24	2.87	-
1187			0.04	0.06	1.38						
•						35 25					

Table 2-1:T3NO vs. BPH: Genes upregulated 5 fold or higher in at least one tumor sample or 3 fold or higher in at least 2 tumor samples.

	MAGE	Gen Bank			Fold/AveTumor/Av i	Cold/MPM120/A	Fold(MPM121/A	Fold(MPM124/A	Fold(MPM122/A	Fold(MPM123/A !	Fold(MPM126/A	s of Tumor
	Clone 10	Accession	AveBPH	AveTumor	eBPH) veBPH) veBPH) veBPH) veBPH) veBPH) Samples	VeBPH)	veBPH)	VeBPH)	VeBPH)	veBPH)	V*BPH)	Samples
4488	878469	A A B 70200	40.4	77.	137	0.74	0.52	0.53	0.42	5.54	0.48	-
	4,000,00	44875053	3 3	; ;		6.50	0.78	0.26	0.58	0.07	0.07	-
80 7	0077441	2020/000	t :			0.60	0.35	0.58	0.75	0.11	5.71	-
2 3	244000	//987LW	9 5	- t	3 6	3 6	0.48	5.19	0.72	0.68	0.46	-
181	102801	AAOABASU	9.0	) S. C	 	0.02	5.81	4.0	0.16	0.27	0.91	-
721	01174	27.000	) e	6 6		, s	0.28	0.18	0.76	0.18	0.50	-
2 ;	<b>10776</b>	017674	3 ;	5 5	÷ 5	6,0	770	58	0.41	5.08	0.82	<b>~</b>
1	5	ASSESS	Ę (	3 3	<u> </u>	5 to 0	0.45	0.28	5.73	0.29	0,45	••
1185	307337	N95226	2.56	<b>3</b>	7 :	e e	0.45 7.85	0.0	649	0.47	5.44	-
90 ;	377286	A055491	) .	7 6	9 3	67.0	9.43	7	0.12	0.12	0.33	-
) (i	277786	A4160670	0.08 0.08	5 6	<u> </u>	7 7	9 20	9.14	0.61	0.38	5.28	-
2 2	503545	W38343 AA131518	0.09	0.10	91.1	0.12	0,12	0.80	0.67	0.12	5.27	-
				;								

36

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

FORGILIPHIZZA FORGINPHIZZIA FOLGINPHIZZIA GOTTURON VOBPH) Samples 8.48 3.05 4.01 2.02 1.75 10.23 3.22 17.00 12.53 Fold (upungtha Fold (upungka 20.64 41.52 41.34 41.34 11.84 11.84 16.61 4.04 0.04 0.05 0.05 0.08 0.08 Fold[AveTumoria Fold(AIPM120/A 22.10 lesst 2 tumor samples or 3 fold or higher in at least 3 tumor samples. 194.72 13.52 1.82 158.73 prostate prostate Gen Bank Accession Number 172543 NTR180 AASO/T84 AAO1176 H48548 H48548 H48588 N42090 AAA32103 N31852 H67712 AA663160 AA663160 AAA54587 AAD04412 AAAS3310 AAA67236 H81849 AA421761 H20747 N91003 R91308 AA609485 T72556 H29966 R83407 R61372 A196979 AA703398 R26397 ¥64846 N55288 N50880 R35051 24.3841 24.2003 267.528 231003 2000 786150 646007 \$500 24.5768 136605 SET 170753 712282 28-1701

Table 2-2: T3NO vs BPH: Genes upregulated 5 fold or higher in at

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

ode.	IMAGE Clone ID	Gen Bank Accession Number	Tiesue Clutifbution	Secreted/Me mbrane Bound	AveBPH	AveTumor	Fold(AveTumor'a Fold(NPM120/A Fold(NPM124/A Fold(NPM122/A Fold(NPM123/A	Fold(MPM120/A veBPH)	Fold(MPM121/A veBPH)	Fold(MPM124/A (	Fold(MPMf22/A /	od(upuizaa F Vebph)	old(NPM125/A veEPH)	of Tumors Samples
ន	825785		Vavo.		55,	8.57	•	9.24	6.98	2.85	6.31	9.04	95° <b>7</b>	•0
\$	49659		r and		20	8,	v	2.31	3.63	6.80	3.61	13.00	3.10	w,
ŧ	417078		•		280	4.38	40	2.70	3.62	3.68	171	8,09	9.64	<b>1</b> 0
7	840751		•	>	ŗ	806	40	3.31	10.29	3.87	2,36	7.56	4,13	10
3	241821		•	¥	8	6.43	<b>u</b> n	3.85	6.61	6.92	90'8	2.70	3,31	•
1	28063		prostate		1.68	6.52	10	4.28	6.83	4.57	7.80	4.62	2,43	₩
\$	124128		•	>	126	6.16	<b>40</b>	2,53	5.00	3,89	4.48	8,48	5.11	100
\$	123862		Activity		4.70	22.13	*0	233	3.01	3.66	11,43	<b>4</b> .82	4.20	10
4	134719		Lopos Color		67,0	04.7	*	5.02	7.59	4.08	. 76'9	3.63	2.83	*0
#	287745		brank		3.81	18.41	**	2.38	4.28	3.76	2.64	3.82	5.12	•
<b>.</b>	271141		•		100	900	•	0.70	8.80	4.19	3.70	6.37	95'7	*
8	1240414		•.		0.79	3.60	*0	1.68	4.13	3,48	7.19	73	4.47	ш.
5	307249		testis		338	15.28	<b>ю</b>	9.25	5,34	07:0	6.39	3.20	3.63	₩.
3	783840		Overy		0.13	0.59	<b>~</b>	4.62	5.10	6.45	4.67	1.71	<b>4</b> .9	₩.
<b>,2</b>	607720		d d		0,32	<b>8</b> 21	•	3.68	3.50	2,94	3.83	6.46	\$.83	•
3	283070		(seg)		8	0.12	•	3.87	9.89	3,48	5.56	7.	4,45	••
3	265005		Ą		900	070	•	6.18	3.28	3.22	9.00	979	6.16	•
8	343987		<b>Pone</b>		972	36.37	•	6.16	7,99	3.20	3.85	97	3.50	•
6	416855		•	>	9,0	1,88	•	2.15	7.06	3.45	3.00	5.14	icc	•••
3	842848		Pund	<b>&gt;</b>	1.15	4.62	-	3,40	4,63	2.64	7.	5.85 5.85	3,05	<b>5</b> 0
2	40690	•	hrafn		900	0.18	•	3.25	3.83	4.50	3.68	4.50	231	<b>4</b> 0
8	359081		prostate		22,22	84.82	•	4.70	95'9	3.16	3.29	3.40	<b>3</b> .	•
5	126230		brain		800	0.10	n	3.10	3,38	3,88	4.66	4.90	0.80	₩.
2	375682		prostate	>	15.66	45.43	•	3.03	3.29	3.51	3.03	3.10	1,45	•
2	1468260			¥	290	22.36	ħ	18.89	1.60	50.34	105.24	0.80	19,00	-
z	743628		egoq	¥	99'0	12.90	ដ	29.79	48.11	29.56	0.86	23.62	1.26	•
\$	1390564		prosteta		18.	33.00	81	49.03	48.43	3,22	1.15	2.77	3,51	•
8	41647		Chard.			7	<b>5</b>	9.65	19.8	<u>*</u>	57.03	15.40	1.72	•
6	741944		prostata		0 <del>.</del>	2,23	2	2.76	4.13	1,26	8.62	48.21	6.35	•
3	נמאמו		prostate	¥	0.13	<b>1</b> 38	F	22.40	22.74	3.29	979	2	1.01	•
3	487165		colon		100	0.09	<b>6</b>	13.49	10.45	13.83	13.97	8.	8.	•
2	1456424		Meed		80.0	Q.69	€	10.16	21.53	1.58	8.31	<b>3</b> .	2.76	•
F	380155				100	0.11	<b>40</b>	10.01	0.73	2.20	24.30	4.57	6.15	•
Ę	144740		prostata		0.10	127	1	7.58	10.38	6.82	10.48	1.84	2,63	•
ß	24915				0.83	6.97	•	3.07	53	437	2.85	10.25	15.67	•
Z	30100			>-	<b>9</b> .70	60.24	•	9.37	12.00	972	6.80	877	373	
78 -	, 21,990		mescle		11,61	70.00	•	9.60	6.06	10.11	2.10	1.74	10.42	•
2	, 035287		•		56.0	5.35	•	8.95	11.77	5.12	5.02	151	2.67	•
-	2		,		100	0.07	•	16.58	53.0	<b>8</b> /4	181	2.05	3.10	-
92	1 175184		•	¥	200	1.68	vo	2.60	6.47	15.0	6.43	6.07	2.7	-
2	, 901109		breest		0.76	3.80	wo	3.93	6.17	7.31	8.02	2.78	2.67	•
8	, 908.187		prostate	>	1.78	7979	<b>w</b> o	6.20	9.84	6.65	05,1	0.28	5.90	•

Table 2-2: T3NO vs. BPH: Gene upregulated 3 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

			į	Secreted/Ne		•		Fold[Ave]umon's Fold[Aphi21A Fold[Aphi24A Fold[Aphi22A Fold[Aphi23A Fold[Aphi23A Fold[Aphi24A Fo	Fold(MPM121/A	FOLKIMPM124/A	FOM(HPM122/A	Fold(MPM123/A veBPH)	Fold(MPM116/A veBPH)	gentles
Oder	INAGE	Accession	Cletribution	mbrane	AvedPH	AveTumor		veBPH)	A48FE	(u. bogan			0 47	. •
						•		\$7.8	6.67	6.33	1.52	3		•
*	754034	AAA79058	•		0.02	0.10	o 1		13.78	4.92	2.80	1.74	376	<b>-</b>
: :	970500	TOTANA	poop		\$	0.18	₩	707	8	200	282	7.0	8	•
3 .		163000	200		. 200	90'0	*0	8			11.80	629	2.97	₹
3	200	900			3.53	17.34	\$5	32	i.	3 3		2.70	10.45	4
3	616262	Nestos			7	2.17	*0	3.19	d e	9.79	3 ;	157	11.65	•
3	296210	N74392				8	•	3.6	0,53	3.	1.14		5.71	•
28	40402	R52089			70'0	8		1.99	4.80	2.74	9.28	7 .		•
18	858153	AA633811			8	•70	•		9.73	7.83	3.55	200	3 !	
\$	704697	AA282253			077	128	n		40.40	**	£28	1.10	7.07	, ,
3 8	A1.00	A A 4 2 0 1 1 B			976	15.42	¥n	S. C.		5	187	87	7.69	•
3					0.02	0.10	<b>1</b> 0	0.48	3	3 5	87.6	1.85	2,40	•
8	3	200			1.70	707	<b>v</b> r	360	10.61	9.0		37	<b>4.68</b>	•
5	200	AA490263			, t	50.70	40	5.52	5.50 5.50	<b>3</b> .	5.	85.	2.87	•
2	260170	N32072			3 !	:	v	4.85	3.66	3.87	200	} :	9.70	-
8	448309	AA777902		ž	747	1		707	2.42	10.26	 6	10.4		•
: 2	276555	M81758		Maybe	9.0	<b>4.35</b>	n	<b>,</b>		3.73	5.40	9.60	Ţ	,
£ :		***************************************			9	4,15	4	1.85		9	135	5.02	1.7	•
3	7000	907704			170	2.07	•	7,5		P. 0		98.6	5.65	<b>~</b>
8	196387	R88915			; ;		4	2.17	5.43	F.7	3		1 97	•
6	428267	AA004628			=	9.50	, -	3.75	3.54	4.23	10.05	8 1	<u>.</u>	. 4
3	391984	A1003625		×	4.42	19.15	•	9	4.37	3.15	17.0	1.81	3	• •
8	607.00	H77854			1.82	8.25	•	e (	8	2.50	3.26	4.87	2.90	•
2				>	15,01	62.10	•	5.48			1.43	81.8	1.26	•
8				•	0.75	308	•	1.97	3.25	•	2 5	1.75	97'7	₹
ē	2102012	HEESTO		-	2	0.10	•	6.87	320	9.8 L	3 ;	# 3B	2.07	•
ā	800456	AA443090			70:0	2 5	•	2.86	86.0	3.63	4:72	3	7	•
ā	361659	W96189		3	6.37	\$0.00 1		89 1	2,48	4.06	2.17	5		• -
Ş	148225	H13688		<b>&gt;</b> -	7.7	101		- C	321	2.7	10.52	990	07.5	, -
Ş	417867	W90128			13.58	27.72	•	900		8	5,13	8.53	2.31	•
3 3	7,083,1	******		Š	1,	5.81	•	1.27	5 5	9 .	274	2.97	3,41	₹
8	3				1.74	75.9	◄	g.11	20.0	ָּבָּי <sup>ָ</sup>	9	8.17	1.75	•
Ď	8K181					238	4	208	8	);*		177	3.74	•
효	1667833			Š	3 3		7	9.05	475	<b>7</b> .	ř		77.	•
ş	144747				3	Ž	• •	6.10	5.74	3.61	2.10	<b>7</b>		
5	321384				Ŗ	4.70	•	75.0	4.47	8.10	3.29	10.7		, -
Ξ	7,5285				900	0.18	•		23.6	3.70	6.85	1.28	3.	•
: :	256464				0.28	101	•	3 t		5.5	3.38	9.7	3	•
2	266692	N25920			181	879	•	37.		3.82	77	4.10	6.85	•
	0	1			0.83	342	•	<b>*</b>	}		7.7	38.6	0,49	•
= ;		2000	, ş		3.58	13.06	4	4.67	0.01	3	8.17	4.30	6.1	•
2				,	5	3.54	4	282	5.13	900		21.12	2.83	•
‡	25.00	AA630104		-		5	4	4.16	2.79	3.62	90.6		11	•
117	24696	H62061			2	90.00		3.55	5.47	3.61	8. 78 8. 78	;		•
118	7.08	AAA27628			10,52	20.5	, ,		6.29	3.1		376	DA :	, -
2	482010	AAD45709			860	3.54	•	2 3		5.01	9.₩	4.57	178	•
: 1	4.24	CROCOL		ž	27	40.58	₹	5	5 5	¥	2.19	2.78	24.	4
3					8	0.10	•	5.81	3	2 3	\$	5.82	1.66	•
₹.	99	162842			900	33	•	2.09	T.		***			
ħ	653012	AA864104		<b>S</b>	j			•						
								••						•

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Order	LIKAGE Clone ID	Gen Benk Accession Number	Tissue Distribution	Secreted/Me mbrase Bound	AveBPH	AvaTumor	Fold(AveTumorfa Fold(MPM120/A Fold(MPM121/A Fold(MPM121/A Fold(MPM132/A Fold(MPM132/A # of Tumors veBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH) veBPH) Samples	Fold(MPM120/A veBPH)	Fold(MPM121/A	FOM(MPM124/A F VOBPH)	Fold(MPM122/A veEPh)	Fold(MPM123/A F veBPH)	-old(MPM125/A : veBPH)	Samples
	811914	AA454986	•	W	288	10.39	•	3.92	6.10	3.87	3.79	2.79	1,62	~
		147808	ş Q		970	1,25	m	5.38	3.04	3.53	2.17	242	<b>**</b>	- •
128		AA704650		>	9.01	31.38	60	7.44	3.12	4,85	£.	3,48	6.29	•
		AA757401	pooq		220	7.68	n	2.20	5.31	3.00	4.08	4.16	2.06	•
121		R63253	•	>	<del>7.</del>	4.88		1.74	4.81	3.90	3.23	4.16	2.05	•
		N63609	•	>	19.65	88.00	₽,	3.37	5.00	1.36	1.50	6.00	3.20	•
_		AA677461	•	**	990	1.89		1.61	4.23	4.13	4.28	4.83	87	•
Ş		HECCOSE	testia		27	20	•	5.12	6.73	3,43	3.25	0.82	0.85	•
Đ		H48502	embyo		0.16	30	67	3.02	4.77	304	2.28	<b>4.65</b>	252	•
ŧ		AA758154	Color		99:0	174	•	1.87	6.62	3.24	4.60	09.1	3.25	· <b>~</b>
\$		H84095	color	>	76:0	3,25	•	2.04	3.28	4.07	1.86	4.90	3.08	•
¥		<b>FC36364</b>	•		0.38	124	•	1.98	3.01	5.86	3.28	1,47	3.16	•
\$		AA883504	Phymus	¥	1.18	3.84	•	1.87	3.82	4.55	3,37	3.97	7.0	•
_		AA835560	prostate	ş	1.17	3.77	•	3.58	5.86	3.20	0.90	4.48	1,12	•
		AA116537	bread		75.0	1,72	•	1.82	3.32	2.48	4.38	3.33	3.88	•
	744001	AA629009	<b>2</b>	ş	3.87	12.24	n	1.80	3.05	3.27	4.18	611	1.80	•
Ş		N63648	•		3.38	10.70	n	2.38	3.84	3.44	3.88	3.55	181	*
<del>2</del>		AA429602	prostate	Maybe	1.17	3.67	•	3.28	2.51	3.73	2.89	322	321	•
<b>7</b>		AA877208	•		1.02	3.16	6	2.09	3.15	1.69	3.87	3.10	4.59	•
\$		AA708370	•	¥3	0.30	3.1	n	4.28	2.84	4.05	3.20	1.06	1.87	<b>~</b>
3		R44132	Prafr		953	3.	•	502	3.6	3.70	3,14	3.71	<b>34</b> 1	•
Ī		AA400365	prostate		8	000	n	333	3.10	2.05	4.16	0.88	4.56	•
2.		810018	5	>	1.14	3,4	•	2.11	2.81	3.08	3.17	3.53	3,40	•
₹		AA007370	bress		1.28	3.63	6	3.22	3,20	1.89	3,35	27.	3,6	•
147		F36465	•		0.47	137	•	3.15	2.95	3.25	3.12	1.85	or o	•
<del>2</del>		AA700604			42.83	123.16	n	3.86	5.22	3.12	12	0.47	3.55	•
3		44498349	<b>1</b>		0.11	970		3.57	4.62	3.03	1.81	0.36	86	•
₹ 		N32514			<b>6</b> .0	27	8	3,34	1.13	1.13	14.34	134,44	1.13	•
<b>\$</b>		AAA31721	prostate	>	83	3.10	ā	64.02	1.22	0.11	0.61	3,63	22.03	•
₹ 		W16659	¥0,		900	98.0	*	1.39	3.76	0.97	71.38	3.78	2.08	
3		N9411	spleen		880	12.90	ŧ	0.73	1.52	4.77	20.47	44.85	<b>307</b>	•
		N47717	endothellai		820	8	tī	43.16	25.22	6.01	222	030	12	•
		AA115761	embryo		ğ	9,46	t	46.34	4.67	1.83	0.27	18.31	. 280	•
		AA443147	<b>t</b> og		800	0.92	F	0.12	0.12	27	65.00	18.3	3,44	•
		NS9156		>	1970	<b>5</b>	ŧ	20.89	31.83	7.51	1.25	980	1.67	•
		N48259	prostate		8.	10,34	đ	21.89	29,40	4.51	1,88	1.07	228	•
		AM16627	Dress(		<b>4.08</b>	33,07		14.84	26.99	1.81	1.28	0.78	421	•
	_	AA0238987	prostate		2.85	20.02	•	1.07	4:78	50.	5.39	32.28	268	•
	-	AA000851	Pe es		0.07	757	•	3.64	3.42	1.60	<del>1</del> ,38	34.08	273	••
<u>\$</u>		N40577			0.13	76.0	7	0.87	3.83	1.15	32.30	5.68	0.12	••
	-	<b>WA37241</b>		¥N	0.47	3.37	4	4.67	30.78	1.50	3.88	0.00	۲. پر	•
ž	-	AA040861	musche	≨	8	35.89	-	241	478	2.60	21.50	878	1.50	•

Table 2-2 T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Order         Clone 1D         Accression         Distribution         Rob           165         258417         Number         Distribution         Rob           166         26818         Accression         Bibribation           166         185188         Accression         Bibribation           166         185188         Accression         Bibribation           170         172040         Accression         Bibribation           171         172040         Accression         Bibribation           172         172040         Accression         Bibribation           173         170040         Accression         Bibribation           174         172040         Accression         Bibribation           175         172040 <td< th=""><th></th><th>SecretadiMe</th><th></th><th></th><th>Fold(AveTumor/A</th><th>Fold(MPM120/A</th><th>Fold(avetumora Fold(MPM120A Fold(MPM121A Fold(MPM124A Fold(MPM122A Fold(MPM123A Fold(MPM120A Fold(MPM120A Fold(MPM121A Fold(MPM121A Fold(MPM120A Fol</th><th>old(MPM124/A</th><th>Fold(uPut122/A</th><th>Fold(MPM123JA 1</th><th>Fold(UPM125/A veBPM)</th><th>s of Tumora</th></td<>		SecretadiMe			Fold(AveTumor/A	Fold(MPM120/A	Fold(avetumora Fold(MPM120A Fold(MPM121A Fold(MPM124A Fold(MPM122A Fold(MPM123A Fold(MPM120A Fold(MPM120A Fold(MPM121A Fold(MPM121A Fold(MPM120A Fol	old(MPM124/A	Fold(uPut122/A	Fold(MPM123JA 1	Fold(UPM125/A veBPM)	s of Tumora
298417 NG4131 colon 666218 AAA23408 skin 1597389 AAA23408 skin 1757389 AAA23510 hypophalamus 112960 AA235270 hypophalamus 112960 AA235270 hypophalamus 112960 AA235270 hypophalamus 122443 WN8423 schring 247540 MA25426 bone 172308 AA42542 bone 172308 R19876 bone 172308 R19876 bone 172308 R19876 bone 172308 H19876 bone 172308 H19876 bone 172308 H19876 bone 172308 MA25420 bone 172308 H19876 bone 172308 MA25420 bone 172308 MA25420 bone 172308 MA25420 bone 172308 MA25420 bone 172309 MA25420 bone 172309 AA425070 heart 162328 MA25420 bone 1220123 AA425220 bone 1220123 AA445220 bone 1220123 AA445220 bone 172307 AA445220		mbrane	AveBPH	AveTumor	veBPH)	v*BPH)	<b>∨•BPH</b>	(K-18)			761	•
256117		Bound	:	•	•	13.15	16.83	1.72	<b>1.</b>	•		•
150789 AAA37142 - 150163			80.0	97.0		ā	867	428	20.28	270	0.61	<b>3</b> (
1901688 R822942 - 1901688 R822942 - 1901688 R822942 - 190169			3	270	•		2	0.65	4.76	95'0	27.68	n (
1951655 R822842 647647 AA235370 hypothalamus 712360 AA282273 Lood 504761 AA44474 bons 722443 WY16424 bons 727566 WA32645 bons 773264 AA43645 bons 773264 AA43643 bons 773264 AA43643 bons 773267 AA43643 bons 773267 AA43643 bons 726537 AA43643 bons 726537 AA43643 bons 726537 AA43643 bons 726537 AA43643 bons 726538 AA44300 bons 7266154 AA43200 bons 726156 AA43203 bons 726157 AA43230 bons 726158 AA43203 bons 726158 AA43203 bons 726159 AA43203 bons 726169 AA43203 bons			0.02	6.1	<b>1</b> 00 1	2 .	1 5	2.16	16.65	4.00	47	n (
647647 AAA25370 hypothalamus 112850 AA28273 hypothalamus 252443 WY18424 sefermal 252145 WY18424 sefermal 252145 WY18424 sefermal 252145 WY18424 sefermal 277350 AA425650 skin 247350 AA425650 skin 247350 AA425650 sefermal 247350 AA425620 bone 245243 W57424 benesit 245243 W57424 benesit 252543 W574243 benesit 252543 W574253 benesit 252543 W574233 benesit 252543 W574233 benesit 252543 W574233 benesit 252543 W5742333 benesit 252543 W5743333  benesit 252543 W5743333 benesit 252543 W5743333 benesit 252543 W57433333 benesit 252543 W57433333 benesit 252543 W5743333 benesit 252543 W5743333 benesit 252543 benesi			897	3,32	•	<u> </u>	\$ <b>9</b>	100	131	1.05	<b>87</b>	n
2001011 AAA14073	_		0.81	4.56	•	9 1	20.5	20.8	0.72	0.83	. 2.12	n
204101 AAA467725 bone 32010 R45118 bone 32010 R45118 bone 32010 R45118 bone 32010 AA42660 sidn 320105 AA42660 sidn 320105 AA42660 sidn 320105 AA42660 sidn 320105 AA42660 bone 320105 AA42660 bone 320107 AA42682 bone 320107 AA42682 bone 320107 AA42693 brain 61029 AA42643 brain 620208 AA43043 brain 620208 AA46303 brain 620108 AA46303 bone			5.83	32.98	•	E.8.		22	0.15	12.57	0.10	•
222423 W16424 edremal 35010 322423 W16424 isstill 722423 M444244 isstill 72252 M444244 isstill 72206 A4442660 edremal 340266 A4461 547526 A446164 isstill 72206  A446164 isstill 722626 isstill 722626 isstill 722062  isstill 7220			90'0	0.43	vs.	6.28	16:0		20.36	0.89	2972	n
25212 VY10212 STREET ST			670	ū	50	3.13	8.	0.73	38.	11,48	6.88	n
35010 R45118  73257 AAA4844 isrtis 77350 AA32650 shin 36465 AA03745 bresst 24736 A403745 bresst 24736 A403745 bresst 77306 A403765 bresst 77306 A403763 bresst 77306 A403763 bresst 77306 A403763 bresst 77306 A403763 bresst 77306 A40377 bresst 77307 bresst 77307 A40377 bresst 77307 bresst			1	3.31	10	27.2	3.	2.60	3 1	¥7.2	8.28	•
77356 N5702 bore 77356 N43762 bore 77356 N43762 bore 34685 A44366 buse 34886 R19876 bore 34886 R19876 bore 34886 R19876 bore 77356 A443692 bore 77357 N42643 brain 61920 H19446 brain 62634 N42643 brain 62637 A443697 brain 62637 A443670 brain 62638 A443700 bore 726156 A443700 bore 726169 A443700 bore			3 5		<b>V</b> 0	2.52	2.75	808	800	ŧ ;	2	n
277309 NST002 bone 24665 AA42660 tkin 246665 AA42640 tkin 346865 AA42640 bessit 247326 T81839 bone 377326 AA42692 bone 277327 NST463 bone 277327 NST463 bone 277327 NST463 bone 276541 NST643 bone 276541 NST664 bone 276541 NST664 bone 276541 NST664 bone 276541 NST664 bone 276541 NST666 bone 276641 N			7.	3 7	, w	7,68	12.12	2.47	80	2 6		•
771208 AA42660 skin 344665 skin 344665 skin 34665 skin		<b>&gt;</b>	<b>8</b>	<b>S</b>	s ve	1.82	1.80	<b>3</b> .	12.13	2.89	3 5	•
247.547 N.26136 brest 1 247.547 N.26136 brest 1 772396 A.4425632 brest 1 770286 A.4425642 brest 1 51870 H19446 brest 1 2263-74 N.67439 brest 1 2263-74 N.67439 brest 1 2263-75 A.412647 brest 1 2263-75 A.412647 brest 1 2262-75 A.445730 brest 1 2261-75 A.445730 brest 1 2261			Ç. ;	BY / 8		1.52	124	13,33	900	7	3 3	
247547 N.95136 bone 24828 R.19876 . 77226 A.425922 bone 77727 M.42592 bone 77726069 A.442592 bone 77726069 A.442592 bone 77726069 A.442592 bone 77257 W.02426 brain 770269 A.442592 bone 726524 W.66139 brain 770269 A.442592 brain 77252 A.442592 brain 77252 A.442592 brain 77252 A.442593 brain 77252 A.442593 brain 77252 A.442593 brain 77252 A.442593 bone 77257 A.442693 bone 77257 A.442693 bone 77257 A.442693 bone 77257 A.442693 A.443083 bone			3	2.00	<b>.</b>	3	14.29	1.01	4.85	a e	100	•
4489.6         R18876         -           77329.6         731539         piacanta           77320.4         AA43692         bone           277327         N45463         bruin           81820         AA43692         bone           26343         N65436         bruin           26343         N6556         -           26344         N6556         -           26567         AA12843         brain           26567         AA12847         brain           26587         AA12843         prainis           26587         AA13843         prainis           26587         AA48570         king           26587         AA485373         brain           264624         NIST86         prainis           264627         AA487305         brain           264628         AA445300         breat           264678         AA467303         breat           724678         AA467303         breat           724678         AA467300         breat           724678         AA467300         breat           724678         AA467300         breat           724678         AA467300 <th></th> <th></th> <th>9</th> <th>90.0</th> <th>•</th> <th>2 3</th> <th></th> <th>A.74</th> <th>24</th> <th>1,68</th> <th>25</th> <th>n</th>			9	90.0	•	2 3		A.74	24	1,68	25	n
72296 75559 plecenta 773204 AA42662 bone 277327 N57429 brain 61620 H19446 brain 61620 AA420646 brain 61620 H19446 - 28543 N52543 N52549 brain 622557 AA126407 brain 622557 AA126407 brain 622557 AA126407 brain 622558 N89499 - 285624 N57549 brain 623547 AA45370 brain 643547 AA45370 brain 643545 AA45350 brain 72307 AA45350 brain 623547 AA45350 brain 643578 AA45358 brain 643578 AA45358 brain 643578 AA45358 brain 643578 b			0.14	0.67	<b>1</b> 0	B (0	3	57	137	4,55	0.13	•
773204 AA42892 bone 777320 AA42892 bone 777028 AA430448 bruin 81820 H18448 - 226343 N2543 bruin 62635 AA13640 bruin 62636 AA13640 bruin 62636 AA13640 bruin 62637 AA13640 bruin 62638 AA45470 kung 62638 AA45470 kung 62638 AA45373 bruin 641507 AA67348 bruin 641507 AA67348 bruin 641507 AA67348 bruin 682467 AA45373 bruin 682467 AA45330 bruin 72458 AA45330 bruin			89	970	<b>u</b> o	) (P)	\$ 9		43.00	2,45	3,58	•
777228 M57483 bruin 770288 briin 770288 briin 81820 hriin 81820 briin 81820 hriin 81820 hriin 81820 hriin 828543 hriin 828543 hriin 828543 hriin 828543 hriin 828543 hriin 828543 hriin 818245 hriin 818245 hriin 828543 hriin 818245 hriin 818245 hriin 8182457 hriin 818245 hriin 828457 hriin 828454 hriin 82			12.74	<b>86.08</b>	<b>5</b> 0	1.47	<b>3</b> 1		87.8	1.05	7,68	77
770088 AA430445 brain 61620 H19448			g	600	•	1.87	170	3 9	2	0.26	87.	n
770026 AAA30055 Grann 770026 AA30055 Grann 770026 AA30055 Grann 72637 A 126427 breard 806436 AA43005 Grann 72647 breard 806436 AA43005 breard 806436 AA43005 breard 806437 AA487375 breard 406737 AA67365 breard 725977 AA673657 breard 725977 breard 725977 AA673657 breard 725977 breard 725977 AA673657 breard 725977  breard 72597			7.38	35.36	10	7.28	8.45	G. 16		243	86.0	•
2645.4 H194.46 265.54 H194.56 226.57 H126.5.6 226.57 H126.5.6 226.52 H457.54 226.52 H854.97 226.52 H864.97 226.			Ę	0.12	•	÷.	8.13	7.	3	3	183	•
286343 N25553 N25554 N25554 N25555  N25555 N255555 N25555 N255555 N25555			5 5	01.6	•	9.62	9.86	3	<b>5</b>		121	n
226577 M61284 Prain 526267 A4128407 breast 526267 A4128407 kung 226268 M69499 226268 H69499 226268 H69499 226268 H69499 226268 H69499 22627 A413253 proteile 67367 748225 heast 489637 AA45745 breast 489647 AA45745 breast 1469789 AA445700 heart 724589 AA44520 breast 724589 AA44520 breast 724589 AA44520 breast 724589 AA44525			2 !		•	8.4	12.80	<b>8</b> .34	0.0	Br.	į (	
22556 AA128-407 breard 60528 AA458-710 kinng 20259 AA458-710 kinng 6178-7 T4822 breard 48867 AA459-710 breard 48867 AA459-710 breard 48867 AA459-710 breard 723015 AA459-710 breard 723015 AA459-710 breard 723015 AA459-710 breard 723015 AA459-710 breard 723016 AA459-710 breard 723016 AA459-710 breard 723016 AA49-710 breard 72			2	1.87	• •	280	7.96	1.80	623	174	3 1	• •
806256 AAA128407 bresst 806258 AAA4570 king 222853 N05489 - 205853 N05489 - 205852 N05489 - 205852 N05489 - 205852 N05489 - 205852 N054853 Presiste 617397 T46325 Presiste 224624 N054789 Presiste 224624 N454789 Presiste 224624 AA43040 Presiste 224624 AA4304 Pr			8	22	• •	5	1,67	96.0	9.74	7	1978	, ,
202556 AAA54570 king 202556 N85469 - 2 205628 H57545 Frosisis 67397 T49225 Issibs 811022 AA455373 bread 459637 AA419178 prosisis 24457 AA47745 king 823497 AA47745 king 823497 AA47345 bore 230155 AA443300 heed 723972 AA43300 724578 AA43303 bore 826174 AA43045 bore 826174 AA43045 bore 826174 AA43045 bore 826174 AA43045 bore			0.02	0.07	•	36.00	0.18	0.28	2.13	<b>9</b> 6.	E 1	, ,
203007 N73563 - 203007 N73563 - 203007 N73563 - 203008 H57365 Prostale 617023 AA45577 Prostale 220425 N847865 Prostale 620467 AA457305 Prostale 220425 AA457305 Prostale 220425 AA457305 Prostale 723072 AA45732 Centric 723072 AA45732 Centric 723072 AA457307 Prostale 723072 AA45732 Centric 723072 AA457325 Centric 723072 AA45732 Centric 723072 AA45730857 AA45732 Centric 723072 AA4573085 AA45732 Centric 723072 AA45732 AA45732 Centric 723072 AA45732 AA45			90.0	0.28	•	2 4	3	990	3.72	15.21	3.80	n •
205028 H57565 Fratise 67367 T49225 Issibs 811028 AA45377 breast 499037 AA101878 proutsie 224424 N64799 breast 499037 AA457305 breast 4903188 AA45300 heart 72397 AA45300 heart 72397 AA45230 breast 829174 AA502543 bore 829174 AA502543 bore 596035 AA10087 Neuron 596035 AA10087 Neuron			900	27	<b>~</b>	0 6	97.0	79.6	2.18	1.01	171	n ·
705628 H57565 preside 61397 T46325 issib 811028 AA45537 bress 496937 AA41978 - 841507 AA4799 - 841507 AA6798 iurg 82497 AA6799 bore 220155 M47015 - 1469788 AA4300 heed 724972 AA4300 724598 AA43097 bore 846174 AA50630 bore 846174 AA50630 bore 846174 AA50630 bore			0.42	1.8	•	75.1		15	1.28	1,92	6.70	n
61307 T (48225 18886 811023 AA45537 break 495037 AA101878 presiste 224624 N64789 - 841507 AA48739 lung 822407 AA68739 lung 822407 AA68730 hung 72407 AA43200 head 72407 AA69260 bone 724598 AA43200 head 724598 AA473087 Neuron			0.51	2.18	<b>~</b>	99.7	<b>3</b> 5	890	08:0	8970	9.45	n
811023 AAA5377 bress 49637 AA101878 - 224624 N64789 - 841307 AA48738 lung 832497 AA48739 lung 8320155 N47012 - 1469188 AA83733 - 724589 AA44300 heed 723972 AA83933 bone 848174 AA59630 bone 425168 AA47522 certik 425168 AA47523 heed			0.01	90.0	<b>~</b>	97.9		89.0	7.56	1.37	8.82	n
459637 AA101878 proutale 234624 N64796 - 641507 AA457365 lung 632497 AA457365 lung 1468788 AA45730 heed 723972 AA45230 heed 723972 AA45230 heed 723973 AA45230 heed 723973 AA45230 heed 723974 AA50826 centre 986874 AA10837 Heed			8.3	20.42	•	6.5	9	17.	8.01	744	1.30	n
29,427 AA,6736 'wg 82,497 AA,6736 hwg 82,497 AA,8736 hwg 230,155 NA,7025 'wg 74,598 AA,4300 heef 72,5972 AA,8233 bone 82,6174 AA,50362 centh 42,6166 AA,01970 bone 56,6256 AA,30857 Neuron		ž	288	11.62	•	3.07	2 4	979	777	6.69	.75	•
641907 AA487385 Iurg 622497 AA78604 bone 220155 IA7012 1468788 AA83735 *** 724878 AA43283 bone 826174 AA50620 722578 AA47522 centh 425169 AA470857 Neuron			4.58	97.9	<b>-</b>	80.2	2,50	01.4	379	1.15	2.08	-
823497 AA578604 bone 220155 NA7012 1468788 AA833735 *** 724878 AA43263 bone 826174 AA590830 732578 AA47522 centric 426169 AA470837 Neuron		>	ğ	4,28	•	10.96	763	0.28	9.36	a.re	ž	n
280155 NAT012 1409788 AA83735 • 1409788 AAA4300 heed 723972 AA82683 bone 828174 AA82683 bone 428189 AA44521 centh 428189 AA401870 bone 586826 AA130857 Neuron			500	0.14	•	0.40	2 6	14	2.63	6.9	8.	n
1469189 AA43333 heart 723972 AA432593 bone 826174 AA450520 teart 723978 AA41522 teart 428169 AA41593 bone 566256 AA130857 Neuron			90.0	0.25	•	2.52	A7-7		18	89.1	17.	•
1808100 AA000100 721972 AA402500 bone 826174 AA50620 bone 182515 AA40522 certit 478166 AA01870 bone 586826 AA130857 Neuron		A/A	0.48	1.79	4	2.93	4.	B7'0	5	58.7	11.30	n
721972 AA202530 bone 829174 AA202530 bone 723576 AA47522 canti 425159 AA47522 Canti 526825 AA130857 Neuron		į	800	0.10	•	3.05	1.55	\$	3 8	22.0	0.74	•
826174 AASOG20 732516 AAA17522 centin 426186 AAO1870 bone 986826 AA130857 Neuron			00	8	•	3.91	7.78	<b>3</b>	5 6	12	8.10	••
836174 AAS0620 732516 AA447522 centi: 426186 AA001870 bone 986826 AA130857 Neuron				ž	•	6.41	6.16 6.10	96.0	er o	} ;	:	•
782575 AAA47522 cerdit 428166 AA001870 bone 586826 AA130857 Neuron			3 3	; ;		8.55	7.01	2.57	9.68	9870		
425166 AA001870 bone . 586826 AA130857 Neuron			۶ : 5 :	2	, ,	200	3.10	4.57	10.12	<b>3</b>	0/1	, ,
586826 AA130857 Neuron			877	6.77		2.78	2.15	2.18	8.17	8.4	9.79	,
		¥N.	1.17	4.49	•	i u	i					

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

ò	IMAGE	Gen Bank Accession	Theue	Secreted/He mbrane	AveBPH	AveTumor	Fold(AveTumor/A	Fold[AveTumoria Fold[Arph12]A Fold[Arph12]A Fold[Arph12]A Fold[Arph12]A Fold[Arph12]A & of Tumors 	Fold(MPM121/A	Fold(MPM124A F	Cold(MPM122/A F	OLIMPHIZZIA F VOBPH)	old(NPM12S/A	# of Tumors Samples
	_	Number		Bound			(u.da.					•		•
707		AA417012	testis		20'0	90.0	₹	7.50	9	0.40	2.80	9		•
ž		44450578	Į.	>	7.85	29.76	•	3.76	58.G	1.10	2.48	20	2.62	- n (
2		AA133959	Ş	•	0.12	70	•	1.62	0.0	1.78	6.89	5.74	6.57	יני
1 5		LASTARA	a procedu		080	33	•	5.28	8.02	27.23	1.69	0.76	1.61	<b>n</b> (
? ;		A A DOTT BA1			0.11	0.43	•	3.08	1.05	3.00	3.68	15.0	¥ 00	n (
; ;			-		18	0	•	0.50	0.29	70	4.24	12.61	S;*	<b>r</b> > (
2 3		2000	The state of the s	1		2 2	•	1.02	1.89	3.14	6.23	90'9	2.83	n
		AA703248		Š	770			2,83	70'9	2.57	2.67	3.32	85"7	n
Ř		KI I	•		; <del>;</del>	5 5	• •	5.05	2.30	2.70	1.07	4.57	6.14	n
			• }	4.4	} ;	9	• •	2.83	20.0	1.92	4.91	6.33	273	m
2		20/70E		Maybe	3 3	· ·	- ◄	2.41	7.49	3.46	273	3.62	2.52	n
<b>1</b>		AA707219		717	;		• •	3.63	6.4	2.14	9.14	2.01	37. 1	n
912		K90472	<b>1</b> 00	Š	}	5 6	. •	48.8	4.18	2.20	4.29	120	2.78	n
		MOT/16	• }		3 6	2 2	- ◀	30	3.90	1.60	278	7.17	2.81	•
8		A4424048		<b>§</b>	3 8	3	. •	2.95	6.36	Z,	5.77	22	151	n
Ñ		H95976	Oun Oun	Meyoe	5 6	70.01	, ,	g	2.87	2.75	6.21	4.11	1.47	n
<b>8</b>		N33263	•			7	• •	5 6	98	3.48	7.92	<b>3</b> .	1.82	n
R		AA496455	musce		2 3	\$ ;	• •		9	1.82	1.7.	3.01	7.58	•
ã		AA485166	ecoq		2.68	, i	, ,		172	3.77	7.25	2,34	2.03	n
ñ		AA709333		<b>K</b>	<u>}</u>	ξ ;	, e		22.0	797	38	2.26	320	n
ន		AA431064	#DIDE		3	8	, <b>.</b>	2	:	707	625	3.48	2.49	•
Ħ		H29600			3	77	m (	3 5	:		3	22,	205	•
ន		R05810	<b>pun</b>	<b>W</b>	1.82	0.62	<b>n</b>	70.1			9 9	0.40	1,77	•
8		AA434404	adremed		85	27.68	<b>1</b> 0 (	0.20	2 4		203	5.60	121	n
ដ		AA460827	Dress!		<b>3</b>	<b>8</b> . 1	n e	2 2	9 ×	}	4.16	1,12	1.30	n
ā		H25020	prostate	٧»	2:01	7.07	<b>n</b> (	18.7			8	0.0	0.00	n
Ħ		AA176833	Pung		0.0 10.0	3	ומ	7.87	2 5	3 5	3	£1.1	90'9	•
Ħ		AA284285	spleen		0.0	B. (	<b>7</b>	8 1	2 3	2 5	1.0	1.58	6.0	•
ដ		AA703114	•		80	772	n (	BC:0		, se	1 2	. 81.1	ž	-
Ħ		AA778675	Pert		1.17	6.35	n	9	ė į	2 3		E	27	•
ដ	882483	AA676598			2.76	<b>0</b> .30	es (	<b>A</b> 1	987	2.30	5 6	10.17	7	-
ŭ		AA424535	Money		67'0	1.67	<b>.</b>	20.1	• :	- 1	2 ;	\$	**	•
ន		R06479			8.0	0.10	n	2,12	<b>.</b>	27.5	6 6	2	\$10	- 67
ñ		H76328	•		0.07	0.72	en	1,97	35/	<b>.</b>	3 8	į į	8	-
3		AA181179	poue	<b>\$</b>	6.76	22.0	eo	3,63	3,85	2.67	20 60	3 8	8	e er
24		R00284	hambh	¥	2,30	7.72	n	422	6,13	4.42	8 9	2 2		. •
242		WEDGDS	•		0.72	2,40	n	5.76	2.60	5,7	8	2 :	e 6	, ,
2		RODATE	etateoro		ŝ	3.43	0	237	. 4.72	<b>5</b> .88	8		8 ;	
77		HIMIN		>	3.	6.17	n	2.78	70.6	1,50	57.7	3.56	5	rs (
		WADELL	ş	•	287	70.0	•	0.45	808	6.63	2	201	<del>.</del>	n .
? ?			, efetade		8		. <b>e</b> 9	19,1	4.36	6.28	4.13	2.87	5	•
:		1115040		>	170	8	e	202	4.53	3.17	4.45	מז	23 23	•
		44006016		- 🛪	5	98	n	1,98	243	727	84,1	278	77	n
ţ		Average		Ę	į	<u> </u>								

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

				- August			#Turbus_TumodA	Selection of Codes and Cod	Fold(MPM121/A	Fold(MPM124A	Fold(MPM122/A	Fold(MPM123/A F veBPH)	Fold(UPM126/A VeBPH)	serrings
	MAGE	Gen Benk	Tasse	mbrane	AVEBPH	AveTumor	VeBPH)	(NBPH)	VeBPH	(Hd8eA	<b>*</b>		•	-
ě	Clone 1D	Accession	Distribution	Bound					275	3,49	6.75	3.	8 ;	
		NO POR			7.69	25.20	•	86.7		7	7,7	2.25	9. 20.	n 1
548	790Z9Z	H98812	3	;	13.67	57.76	•	2.00	ar.		4.85	1,83	3.72	n
8	741067	AA478438	pane	>	2	4 .	*1	2.84	<u>ə</u>	977		2.82	121	n
52	400178	A121206	rhand V		2		•	2.8	<b>4.</b> 51	77		484	3.07	•
252	208741	H63176		>	8	2 7	, ~	2.01	2.14	2.13	8 6	37	9,02	n
Ş	416411	W86466	8		3.	33.5	, ,	200	16.5	1.93	g :	3 3	\$08	•
3	418002	WRSSA	•		90	1.62	, (		135	3,60	3.		2	n
4		102626			0.85	1.70	ימ	2.5	0.19	0.19	2.85	4.0		•
2	8				9.05	0.17	es	9.6	6	4.15	1.74	<b>8</b> 0:	5.	•
ž	1202	H86518	• •		1.17	3.78	6	6.57		91.	3.80	202	ž	3 1
257	1232	HD6525			, c	0.80	es	2.48	76		17'9	3.80	203	n
<b>393</b>	110767	100639	eje 4		3 5	70	n	1.50	4.12	747	19.	3.17	2.10	n
3	121948	T07782	ar qu		÷ ;	4,0	•	4.87	5.D6	77		2.80	57.7	n
92	762812	AAA48251	brein		8.	2 9		1,80	101	2.67	476	27.6	3.25	n
181	145513	R77718	ptdr		0.67	9 1		230	527	58.	3		980	6
•	415110	AAABBBBB	pone		<b>6.</b>	19.70	•	9	ž	3.47	<b>5.80</b>	27.6		•
Ä			a colore	Maybe	3	4,52	6	<b>8</b> 6	5	3.78	320	237	9 7	. •
ă	B680L2	Hood	Ī		9.30	1.16	6	87	3 :	2.43	2:00	326	281	. •
ž	107007	KENTO	• }	March	3.18	10.13	e	€.88	3 6	260	5.88	1.06	88	, (
<b>3</b> 8	416374	W86202	200	adkam Mayor	000	6.43	၈	2.80	ia i	3	8.08	323	<b>2</b>	n
8	122 123 124 125 125 125 125 125 125 125 125 125 125	AAB27405		Š		8		173	4.27		806	3.77	202	n
267	137017	F35665			3 3	9	**	2.32	3.85	3.5		1,72	757	60
99	267634	N34117			9 3	3 5	· #7	4,18	151	2.43	3 ;	2,2	4.70	•
289	430253	AABV9427					• •	3.45	3.26	त्र	7 7	88.6	2.10	0
E	1475734	AA572692		¥	7.30	9 5		1.48	6.62	335	<u> </u>	3 :	280	**
Ę	268476	9209ZN			5.5	80°1		3.66	4.92	3.86	3	3	4.13	n
E	79032	161899		<b>&gt;</b>	15.17		. "	2.46	15.6	2.85	3	, F.	3.37	n
Ę	771023	AM27978		<b>&gt;</b>	1,12	10.5	, ,	1.06	2.46	1.78	ž i	3	75.7	•
1	811101	AA485978			2.17	6.72	, e	38.	3.31	2.15	121	3	121	•
378	77017	NZTRACE	•		8	5 5		3.50	17.7	3.84	6.73	3	230	•
	448117	AA702480	•	N/A	<b>3</b> 2	6.10		2.08	2:12	331	80'0		445	
Ę	796674		3		0.48	e :		1.85	132	77	926		3.82	r
27.8	45128				6	<u>.</u>	. "	2.21	3.82	4.69	3 5	2.28	3.78	n
27	214624		prostate			9.6	. •	4.38	4.10	1.80	66.1	97.5	1.91	9
285	743323				9	3	•	101	<u>r.</u>	32	ğ i	3	3.69	•
28.	80050	763246		¥2	12.72	8 1	. •	2.12	4.97	1.27	ž	<u> </u>	308	•
282	1500162			ž	 	e .		2.45	3.22	5.41	777		7	•
ž	268234		<b>Dury</b>		1.72	P	, .	2.99	4.10	3.83	zor.		801	•
1	100623		•		5.	ž.	, .	2.41	3.80	2.86	<b>3</b>	} :	1.32	•
ţ ;	36	TANA		KN KN	19,82	60,03	,	90	5.11	2972	<b>4</b>	• ;	! ;	•
3	3000	44469670			<u>5</u>	ĭ	n	3 5	987	2.37	402	3.5	787	•
8		1	,		0.57	1.71	•	11.1	:	321	4.15	4.88	181	
191	418808		•		1.16	3.50	n	1.73		733	3.89	4,83	2.18	,
22	242686				8	90:0	•	2.52	2		22	329	5,43	n
88	364352	AAG22486			8	Q.17	•	1.60	3.79	8	}			•
8	244050	_			į			7						

Table 2-2: T3NO vs. BPH: Gene upragulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

	٠																																							•	,	- 144
Samples		•	•	•	n	n	**	•	•	•	n	•	•	n	n	n	n	•	<b>5</b>	n	n	n	••	•	n	es	n	n	n	17	6	n	•	n	•	n	,	n	n	n		**
Fold(MPL4126/A veBPH)	2,5	29'0	14	270	187	3.19	2.47	3.8	3,5	673	27	<del>1,</del>	<u>;</u>	8.4	1.17	<b>1</b>	I.	1.74	2.40	0.85	2.19	1.53	0 <del>.5</del> 2	3,82	4.47	3,32	27.2	£14	700	3.07	6.08	5 <u>.</u>	3.01	3.86	1.73	4.62	331	15.	372	4.83	\$	930
old(MPM12WA veBPH)	2,68	83	1,07	1,62	3.10	1,63	3.80	0.80	2.04	1.06	<b>3</b>	3,86	4,41	3,43	16.2	8	0.30	<u></u>	233	330	1,62	08.1	2.83	3.64	4.45	3.66	322	3.31	1.07	2.80	0.21	3,56	3.26	0.70	1.62	3.58	3.01	3,72	3,41	0.24	4.79	4.50
id(MPM12ZA F veBPh)	3.80	4.80	5.62	3.26	101	1.80	2.74	2.34	3.08	1.10	120	2.07	5.64	1.57	2.10	123	1.52	1.78	3,52	2.40	3.41	1.24	3.07	1.18	2.89	1.13	3.28	3.46	2.04	0.80	4.52	3,33	1.85	3.99	230	27.0	3.96	7970	232	0.77	300	1.17
Fold[AveTumors Fold[MPM120/A Fold[MPM121/A Fold[MPM121/A Fold[MPM121/A Fold[MPM121/A BolTumors veBPH] veBPH] veBPH] veBPH] veBPH] veBPH] veBPH]	3.39	2.81	1.65	2.81	2.12	:4.83	3.86	1.74	23	3.41	6.49	3.17	1,41	3.29	3.60	5.0	3.73	6.50	3.18	3.04	. 2.50	3.45	3.40	1.38	3.30	2.31	1.01	1.16	323	3.00	1.70	3.72	254	1.83	3.61	4.32	1.14	2.31	Ĩ	3.37	26:0	0.58
[WPM1211A FoI VOBPM]	3.77	3.07	321	3.25	4.11	3.27	3.30	3.38	3.96	7.88	3.14	4.92	3.16	2.11	3.41	3.77	4.75	3.47	324	4.53	324	4.67	4.23	0.09	9970	3.70	3.40	153	3.90	3.47	4.10	202	301	0.62	3.07	122	204	3.70	306	072	9670	77
(MPM120/A Fold veBPH)	1.72	3.30	1.54	4.82	5.58	2.92	1,35	5.57	2.78	3,82	0.99	220	1,34	2,35	3.08	3.67	2,68	3,18	2.20	2.58	3.68	3.73	2.30		0.54	2.13	1.61	1.82	3.69		0.21	1.60	2.03	4.47	3.21	1.16	2.09		1.39			
AveTumon/A Fold( veBPH) v	_	_	_	_	_			_		••	_	••	•		•,	•	•	.,	~	~	e	•	~	4ñ	đ	4	<del>-</del>	<b>-</b>	ที	<b>~</b>	9	<b>₽</b>	79	Ť	e	÷	7	0.62	2,2	3.92	1,17	4.17
	•	•	*,	*	•	•	6	6				**		6	Ð	6	n	60	60	6		C	•	60		6	n	n	6	•	n	n	es	60	60	*	n	n	•	n	63	•
AveTumor	3,15	53.55	870	3	ž	85.4	0.05	3.38	277	2.19	0.14	ā	0.17	40.40	2.50	g	2,86	131	6.17	0.11	<b>4.</b>	1.78	3.58	1.14	900	0.81	2,62	2.08	6.91	0.1 1	0.13	2.28	1.78	4.39	2.27	305	<b>1</b> .5	700	16.82	900	1.89	<b>5</b> 00
Ачеври	8	17,96	0.11	18.18	1,15	<u>7</u>	770	1.15	0.83	0.75	9.05	970	90.0	3.62	78.0	10,0	0.93	97.0	2.10	40	1,50	20.0	13	0.42	0,02	0.30	76.0	0,78	2.18	\$	90'0	78.0	0.88	1.69	78.0	1:1	0.60	20.0	6,57	0.02	0.74	20'0
Secreted/Me mbrane Bound				¥	*	×								Ž	>			>	>		<b>&gt;</b> -						¥					N/A				¥2			¥		ş	
Tiesue Distribution	<b>E</b>		T Can	prostete	brah	•	•	munde				Pe er	d	Poop	prostate	Pone	brath		prostate	bone	instit.				brain		pegpedeun	placents	Ţ.				T .		brest		Neuron	, city	,	her		poue
Gen Benk Accession Number																																										
Close ID	544639	852520	787000	1401252	447418	1476304	FEE	1456807	<b>20103</b>	263769	427800	307244	810843	1480820	26250	366154	282104	344308	<b>45523</b>	196363	267495	757788	22,72	630545	808874 808874	128234	280407	12921	19014	90839	757440	176554	2900	25238 25238	72527	487831	20097	59456	480409	27388	2001	42543
o de la composição de l	2		Ħ	ž			_				ş									92																		- 82	÷ :		2	

Table 2-2: T3NO vs. BPH: Gene upregulated 5 fold or higher in at least 2 tumor samples or 3 fold or higher in at least 3 tumor Samples.

Clone ID	TOTAL POST OF		mbrane	Ave BPH	AveTumor	Weder.	(NABPH)	veBPH}	veBPH)	veBPh)	PORT (MARCH MARCH MARCH) VARIETY (MARCH MARCH MA		
		Distribution	Bound						9	137	1.19	1.85	n
77 2000		j		8	2.65	6	331	3.85	8.50		128	1.79	•
3	W151400	5	1	17.73	47.77	er)	3.17	4.80	9.76	Ę :	1 3	71.6	•
784130	AA432075	prostate	e di di		8	c	3.68	3.87	2.02	1.13	ָרָיָה ,		•
325370	W52208	pod		3 1	3	•	3.07	98.°C	0.85	8.	ř	100	•
288663	N82384	kidney		0.73	<u>S</u>	, «	8	3.62	0.14	3.08	7.02	800	, ,
344359	TTEETW	hing		0.0	0.18	7	2 6		3.34	09.1	1.23	7	n
10003	R16241	•		1.15	2.70	7	3.5		1.82	39.	1.06 60.1	3,56	•
420423	AAOOBUS	•		0.14	0.35	7	3.16	<u>.</u>	, c	4.12	0.85	2.32	•
25.824.9	ATOR55	•		0.62	1.40	~	3.84	27.	3 5	1.52	3.40	3.14	•
166907	207.764	•		600	070	7	0.60	£ ;	3 5	91,	5.01	130	••
ABOCT.	M21380	. }		0.88	2,03	~	1.29	3.24	1.70	- :	8	3.18	n
457088	AA707125			9 5	20.5	7	2.43	3.80	0.18	)e/e	3 5	88	•
665779	AA136870	Dieg (B)		2	2	-	1.20	3.87	0.24	DE E	, i		•
128506	R10154	prostete		5 6	<b>9</b> .0		4.12	3,45	0.42	3.	0.88	200	• •
639048	AA487505	C C		8		. :	720	0.74	56.1	0.74	42.48	37.00	• •
590853	AA158162	endothellal		100	2	2 •	5	0.16	2.38	2.67	3£.78	5.82	• •
50787	H16833	had		90.0	S.	<b>5</b> 0	3 1	1 87	75.0	95.0	1.1	7.7	~
137984	R63085	ekch Fich		670	22	<b>8</b> 0	2 9	e c	97.	6,43	32.14	1.7	~
856585	AA869222	)sand		0,12	ž	<b></b>	<del>2</del>	2 2	2	0.49	97:0	2.53	~
151477	HCZBCIT	£		1.19	8.16	^	7.32	8.83	Ş	3	99.0	1,40	~
84331	180111		>	0.47	3.10	-	15.33		6	1.63	1.01	Ŗ	~
82028	H22802	•		0.25	1.88	7	13.67	8.7	210	0.52	31.38	5.38	~
RKOA27	AARGAADS	rojao		800	650	90	1.07	<b>4.</b>		25.	18.46	16.17	7
43004	******	•		770	<u>\$</u>	60	0.88	<b>.</b>	900	5	<u>\$</u>	20.80	~
	7	,		800	25.0	89	6.98	87	<u>.</u>	r.	11.30	12.03	~
36736	277777	2		828	8.7	wn	707	271	5 8	20.25	2.00	5.30	~
	1	S AN		80	0.16	10	0.47	1.61	<b>S</b>	3	1970	2,40	~
967.67	7000000	Postsi i		18.25	89.08	40	7.88	Z	2.32	, c	988	2.80	
				0.10	94.0	10	14,30	<b>0.10</b>	55.0	<b>.</b>	230	2.82	84
	10801H	1	ž	9.4	16.28	•	6.72	77	<b>3</b> .68	3 ;	3	0.83	~
2//01/2	AA111080		<b>4 N</b>	0.45	1,95	•	6.41	14.60	R.	•	900	14,84	~
2004	Teans	Ş		97.0	0.71	•	1.17	7.40	7.07	2 5	121	34.	7
				500	070	•	11.10	10.84	0.30		881	8.	~
1				9118	0.72	4	1.22	10.53	8 : 6 :	Ş Ç	0.92	10.57	7
		•		0.02	0.0	•	2.68	8.36	3.6	, i	5	15.08	~
17947	•	•		8	0.11	•	1.17	0.38	5.13	<u> </u>	3 :	\$	*
1001200			5	2	0.57	•	10.73	5.96	28	7	3 3	2	~
10435998	-	endotre:	<b>Š</b> :	40.07	5	•	0.81	2.58	2.98	8,45	8		
46 1699	•	pre-gr	Ş	}		•	1.87	2.07	2.68	ភ	3		•
50692	H17628	<b>S</b> quant		6.0	2 ;	, ,	2.76	527	1.60	11.37	6.71	K 3	• •
342926	AA018980	d		£0'.	8	•		821	77.0	ğ	2	10.51	•
122178	T38615			8.	0.11	•	3 .	9	22	780	25.0	280	••
246808	N53214	۵		<b>9</b> 50	2:06	•	<b>3</b> 1	1	26.	2.08	6.39	8:	~
101846	ראשנונים	•		0.61	124	•	23	0/'0	3 :	280	27.0	ŭ	**
200191	232	•						•		7			

	2047	Gen Bank	-	Secretadine			Fold(AveTumon'A	Fold/MPM120/A	Fold(MPM121/A	Fold(MPM124/A		Folguphizza folginphizza folginphizza #	Fold(upu12KA	
į	Clone io	Accession	Distribution	Parige Parige	AveBPH	AveTumor	F VeBPH) VeBPH) VeBPH)	(HdB+A	(MABAN)	WBPH)		V*BPH)	A-BPH)	Samples
į	8007 COV	1007/007	j		•		•	1.28	2.5		7.62	<b>1</b> 2	8.75	~
		and the second			2 6		. •	, e	108		8	10.16	1,08	~
2	2000	0/000			5 5		• •		180		8.	1,86	2.87	14
E !	BC1010	W05423		:	3 6	2 :	• •		1.16		7.10	5.86	230	~
2	50/0	<b>76/4/80</b>	•	Ş	3 :		•		90.3		20.5	200	38,1	~
2	289025	N59626	Pressi		S.		,	3	940			4,0	9	6
8	<b>ET3</b>	174257	•		0.07		n	50.0	6.10 E		8/7	3	3	
š	796646	AA460115	pund		9.12		•	1.38	0.83		8.52	1,56	202	N
2	<b>4</b> 035	HISTOR	•		0.22		•	0.69	1,60		8.85	7.72	223	~
{ }	484774	Albertone	•	***	57		•7	5.71	7.10		3.1	0.65	<b>5</b>	~
3	107.00	4.000004	•	Ş			•	\$5°	273		3	5.85	1.60	~
\$ ;		A A BOOM 9.	. }	*	•		• •	280	22		5.17	1,80	128	~
3 :	0000	*7087044		Ş	1		• •	2.50	200		20.00	850	0.70	~
8	•	M16/43	S		Š		,		1		•	200	878	8
5	742164	W63778	prostate		3		•	0.80	0.20		2	3 3	3 :	
3	795407	AA463318	5		0.13		~	જ	0.07		5,45	28.0	78.0	•
3	10501	7139996	· proetate		90		7	67.0	12		250	3,65	828	<b>N</b>

5

Table 2-3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 2 tumor samples or 6 fold or higher in at least 3 tumor samples.

#	unor samp	ies or 6 fold or	tumor samples or 6 fold or higher in at least 3 tumor sumples.	et 3 tumor eum			AWAIN FORMANIA PARTIES FORMANIANA	Oldestinger -	F-Id/WPM421/A	Fold(MPM124/A	Fold(MPM122JA	FOLGUPHIZZIA FOLGUPHIZZIA FOLGUPHIZEA	FOR(MPH12EA	Samples
l		Gen Bank		Secreted/Me	į	A Townson	Fold(AveTumor/Av	FOIG(MPM120A	OHE BAN	veBPH)	VeBPIQ	Ass.u		. ,
3	INAGE	Accession	11880	mbrane	Avedor	Average	98 H)					22.54	10.38	•
Ě	Close 10	Number	Clatribution	Bound		1	1	140.63	185.70	7.38	277	9	35.00	•
			9(0)0000		8.	130.47	88.78		24.62	11.47	31.08	3		•
_	081887	2000			238	114,00	47.88	47.60		Z 87	5.26	41.24	3	•
~	646037	AA196979	prostate	;		73.10	30.37	31.27	à i	70.07	18.57	45.42	29.52 57.	•
47	. 284701	N64840	prostate	-			30.08	38.52	11.57	10		. 21.41	63.61	•
	670077	AA703396	prostate	¥	3	5	2.5	75.5	8.80	10.00	. !	÷	13.29	•
		ABCORN	•		200	020	0.77		20.64	2.40	18.57			•
_	247	3			24.06	423.65	17.62			0.28	10.57	5	•	•
•	28,003	N20880	<b>1</b>	;	5	44.38	17.34	29.78	701		9.24	10,23	8,3 <b>0</b>	•
	138605	105051		>-	3		2.0	30.48	7.	6. J		47.00	10.56	•
	2000	7023670	alateno	<b>&gt;</b> -	 64:	27.62		97.0	11.04	11,99	10.01		19.24	•
		07777			5	0.12	(8,1,5)		6.32	8.58	7.12	R.		•
	478753	Necles.	į		207	19.56	0.43	£/.5		104	15.8	12.71	<b>,</b>	•
2	61433	H20747				181	7.87	5.89	8		13 61	19.40	9.01	a
¥	195975	R01398	•		R :	3 8	19 55	15.20	2.45			13.28	1.75	w
: :	207678	M62008			<b>6</b>	8		7.01	24.39	10.24	70		¥	10
2	20,07	0.00	1		67.	13.52	10.47	1.9		11.93	6.34	124	200	u
2	2007			;	4.24	12.57	10.18	7.69	10.51		20.21	3,5	8	
<b>*</b>	1030855	AA621761		-	1	The c	7.65	5.69	<b>4</b> .8	0.72	9	5.75	8,20	en
	1031599	AA609485	•		2	10.7	6	3.24	10.05	<b>9</b>	2 :		11.57	₩
2 :			) American		629	1.07	6		\$	5.38	5.43	3	80.	**
2	1000			<b>V</b> / <b>X</b>	1.91	12.10	6.37	770		8.30	879	8	3	•
Þ	825858	•		٤	2	0.20	6.62	7,23	8 9		105.24	08'0	19,00	ŧ
5	130136	. R20813	•		3	2	34.16	18.89	8.	* 'AC	2	23.62	1.26	<b>→</b>
9	1468260	AA884928		ş	2 1	1	22.20	29.78	48.11	28,50	3 8	15.40	1.72	4
۶	743828	AA834378	•woq	¥2	86.0 0	2 7	73.5	9.65	9.81	<b>7</b> .	3. 6		3.62	•
1	10043				9	Į.	1	68.0	26.16	14,54	17.70	1 1		•
5			•		15.97	184.72	B1.21		9	7.67	13.10	77.		•
Ħ	2017	•		>	17.59	158.73	8.02	10.87	3 9	5	13.97	<u>5</u>	8.	,
ĸ	101923	R83407	Drain	•		800	8.86	13.40	10.45	20.5	4.72	3,65	16,91	•
72	487165	AA045074			5 6	ā	8.70	13.54	16.10	<b>9</b>	2 6	4.07	5,11	•
52	306086	N91003		>	8		£7.8	3.8	10.11	7.7	3 5	ž	2,53	•
*	292217	N79180	<b>B</b> ood		90.0	7 .	9 6	7.58	10.38	0.02	10.00		3.81	•
1	076377		amatete		0.10	1.2.1	3	8	10.82	5.48	10.02	į	.,	•
7		•		<b>4</b> %	0.38	2.38	<b>*</b>	2 6	8	10.11	2.19	1.74		•
R	462400				11.61	70.00	8	90.00		6.12	6.02	1.51	70.7	
8	1056172	-	Dinter of		0.95	5,55	<b>3</b>	8			7.48	3.67	87.8 8	•
R	782367	•			6	0.14	5.84	a. 10	5		6.43	5.07	1.1	•
F	195034					- 188	5.48 84.48	2.60	6.47	i i	85	0.26	8.08	•
H	461641			<b>4</b>	ţ :	78.0	40.83	9.20	78.5	g i	:	3.	71.0	•
Ħ	787938	3 AAA52778	9 prostale	<b>&gt;</b>	0/1		10.5	5.74	79°W	0.33	4	203	17.1	•
7	754054	1 AAA79058			0.0	5 6	, T	7,	6.48	æ:3a	3		7.01	•
; ;	195367		•		0.47	7.07		11.55	50.28	3,85	5			•
3	899868		States		5. 2.	48.10	er.c.		25.22	10.8	111	3	1 5	•
8			•	71	0.70	8.6	12.65	2 3	194	5	0.27	1831	70.0	•
Ħ	28182			•	3.0	0.46	12.37	7.8	3 5	1.86	8.52	48.21	5.35	1
3	490718	•			8	22.42	11.81	2.76	2 *					
8	7484	4 AA6258PC	o prostate			i	•	•						

Table 2-3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 2 tumor samples or 6 fold or higher in at least 3 tumor samples.

	INAAF	Gen Benk	F	Secretedille			ASSESSMENT A PROPERTY OF THE P	400000000000000000000000000000000000000	4/2007/00/00/07/17	2 41707101010777	***************************************	- Altertion		A 00 Thursday
PO	Clone D	Accession Number	Distribution	mbrane Bound	AveBPH	AveTumor	Pola(AVE) UnionAV	veBPH)	HABOA .	veBPH)	veBPh)	veBPH)		Samples
\$	287728		•	>	0.51	5.0	10.69	20.89	31.93	7.51	125	980	1.67	•
÷	10H7		prostate	¥	0.13	1.38	10,50	22.49	23.74	3.20	978	252	1,91	•
<b>~</b>	1456424		bread		800	0.69	8.20	10.16	21.53	1.58	8.31	4.84	2.76	•
3	380155				9	<b>1</b>	8.02	10.01	0.73	220	24.30	157	6.15	•
\$	96160		poug		<u>F</u>	11,25	6.59	7.30	5.16	4.03	15.63	2.66	4,40	•
\$	266211		prostate		0.57	27.	8,48	8.50	3,86	13.77	2.12	4.38	6,18	•
\$	693185			>-	9.70	60.24	6.21	9.37	12.99	977	6.80	238	3.29	•
41	428816				0.10	08.0	6.03	8,54	8,44	9.60	4.79	4.07	5.75	n
3	811582		<b>poue</b>	<b>&gt;</b>	3.50	21.22	5.92	9.12	6,48	4.71	<b>9</b> 0°	2.60	8,45	n
\$	428434		•		3.0	3,12	5.83	1,13	4.85	£.4	5.71	11.11	7.68	_
8	795729		peypedsun		0.02	0.11	5,58	. 68'9	4.96	0.83	11.63	400	57.6	•
2	712950	AA282273			5.83	32.08	5,56	6.81	14.24	8.62	0.72	0.86	2.12	•
3	825785		Overy		<b>3</b> 2.	8.57	5.51	3.24	9.08	2.05	6.31	9.01	4.56	n
3	50-07-61		bone		90:0	0 <b>,</b>	5.45	6.26	0.51	1.72	0.13	17.97	0.10	•
3	491706		Dressi		97.0	3,90	5.23	3,83	6.17	7.5	. 9.02	2.20	2.67	•
z	241821		•	¥	<b>50</b>	8,43	5.10	3.85	6.91	5.02	808	2.70	331	•
2	782537		tests	-	7.7	8,80	5.01	2,52	2.75	80.8	0.00	2,48	8.28	•
<b>2</b>	277296		Pone	<b>&gt;</b> -	1.46	7.7	4.97	7.68	12.12	2.17	000	1.13	5.84	•
2	124128			>	125	6.16	4.92	2,63	8.00	3.89	<b>4.</b>	878	5.11	•
2	134710		<u>8</u>		0.20	1.40	4.85	5.02	7.58	<b>4.08</b>	<b>3</b> .8	3.53	2,83	•
8	814306		prostate		9.66	45.42	4.71	5.23	10.10	171	5.26	1.16	2.07	•
5	781233				0.02	0.10	4.70	0,48	4.83	6.38	7.81	6.30	7.60	•
3	34888				0.14	79'0	4.62	5.69	9.05	6.74	1.83	1.88	2.85	•
8	260170		prostate		12.65	69.29	4.61	5.52	5.50	9.33	1.01	1,63	<b>€.6</b> 8	•
3	277327		brafin		0.02	80.0	4.56	1.67	120	2.55	8.18	1,05	. 7.68	•
2	307249		teatta		3,36	15.28	4.54	9.25	5.31	0,40	6.39	3.20	3,63	•
8	7,0066		push		7.38	33.36	4.62	7.28	8.45	8.19	15.0	0.28	2.38	•
5	S1620				0.03	0.12	4.40	1.80	8.33	7.29	6.63	2.43	95.0	
2	128267		pegpedau		1.43	0.20	4.38	2.17	5.43	1.71	5 <u>4.</u> 1	3.90	5.65	•
8	278504		brain		0.06	27.0	*5*	0.98	7.96	1.80	5.23	124	8.83	•
g	205628		prostate		0.51	2.18	4.28	7.69	2.40	5.57	1.28	1.92	8.78	•
Ε.	265005		aldin		9.05	0.20	4.23	6.18	3.28	3.22	6.00	95.0	6.16	,
Ľ	67307		teetis		a.01	90'0	4.18	5.28	. 7.99	0.68	0.90	0.68	9,45	· III.
r.	104624		•		<b>8</b> 2.	6.38	4.03	2.09	6.18	6.46	224	6.60	1.52	,
<u>-</u>	182497 A		•uo4		70	0.14	3.84	0.28	7.63	0.28	. 5178	0.79	75	
r r	126174 A			_	89	0.11	3.89	6.41	6.10	0.90	0.35	133	8.10	
2	4 32900		chard	•	0.12	9,44	3.78	1.62	60:0	1.76	6.89	8.74	6.57	
 F	01230	_	placenta	•	99.0	2; 2;	3.78	6.28	5.02	7,33	1,00	0.76	1.01	
2	Y 0957		spleen		0.01	8.0	3.40	1.86	1.13	5.78	6.41	1,13	6.09	
2	70017			•	0.0	0.23	28:52	3.34	1.13	1.13	14.34	134.44	1.13	***
<b>2</b>	100564 A	-	prostate	_	1.84	33.00	18.02	49.03	48.43	322	1.16	2.77	3.51	~ ~~
K <b>5</b>	<b>₹</b>	-	rostate	<b>-</b>	929	3.10	15.42	64.92	122	0.11		3.63	22.02	
							•							٠.

Table 2-3: T3NO vs. BPH: Genes upregulated 10 fold or higher in at least 2 tumor samples or 5 fold or higher in at least 3 tumor samples.

9 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0
105 205 22.88 22.88 22.88 4.21 4.40 6.38 15.87 15.05 1
Fold(MPM122A FOLDMFM12AA FOLDM
Feld(µPuf12/A ve BPh) 28.47 0.74 1.88 1.28 1.64 1.65 1.56
4
Fold(MPM121/A veBPH) 1.52 0.74 28.40 26.89 18.31 22.58 2.31 16.83 0.41 2.77
Pold(MPM120/A veBPH) 0.73 0.74 21.88 14.84 14.84 15.33 13.67 3.07 13.16 0.88 2.04 11.10
Pold(AveTumorlAv Fold(MPM12)A Fold
AveTumor 12.90 0.18 19.34 33.97 3.19 1.86 5.97 0.48 1.39 44.96
AveBPH 0.00 1.00 1.00 0.01 0.02 0.02 0.02 0.02
Meybe
Tissue Distribution spicen spicen spreakto bresst colon colon
Accession Number Number Number Na4111 AN18152 AM18259 AM18251 TE0111 HZ2833 RZ8068 N74131 AA006196 AA439142
IMAGE Clone ID 201579 600853 282089 731200 81331 52018 24018 24018 764358 1380018
£ 4444488888

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

Table 24; Poor vs Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

Order	IMAGE Clone ID	Accession No.	AveGood	AvePoor	Fold(AvePoor/A veGood)	Fold(mpm31/A veGood)	Fold(AvePoor/A Fold(mpm31/A Fold(mpm33/A Fold(mpm34/A # of Tumors veGoot) veGood) veGood) Samples	Fold(mpm33/A veGood)	Fold(mpm34/A veGood)	# of Tumors Samples
Υ-	753076	. AA436479	1,42	42.16	29.71	2.52	57.56	34.98	23.80	ಣ
. 2	124071	R02586	0.53	321	6.08	0.69	16.64	3.34	3.64	n
m	346321	W79647	0.64	3.29	5.17	3.69	7.75	1.51	57.7	6
4	855624	AA664101	24.18	108.55	4.49	4.12	6.82	0.22	6.80	ო
ĸ	42325	R61073	13.03	62,25	4.01	4.90	5.01	4.68	0.57	ဗ
ဖ	287528		2.13	6.32	2.97	4.00	3.29	1.30	3.29	ю
7	756595	AA444051	1.46	3.89	2.67	0.45	3.61	3.01	3.60	m
΄ ထ	452374	AA700876	0.64	19,98	31.26	0.24	62.30	0.37	62.14	77
o	357105	W83315	0.87	8.54	9.81	0.81	25.33	12.26	0.86	7
9	460798	AA708152	1.24	6.95	5.61	0.47	17.80	3.83	0.33	7
7	284601	N76201	0.27	1.50	5.49	0.82	10.19	0.77	10.17	2
7	480487	AA677708	53.51	237.74	4.44	0.02	8.87	0.03	8.85	7
<u> </u>	248412	N58558	98'0	3.63	4.25	0.95	7.66	0.75	7.64	N
4	212188	H68648	0.56	2.37	4.23	0.93	6.76	2,47	6.75	8
15	782513	AA432030	4.08	16.09	3.95	1.06	1.58	6.19	6.98	8
16	782513	AA432030	2.03	7.78	3.83	1.37	1.28	6.00	6.70	7
11	129128	R10935	0.38	1.47	3.81	0.92	3.71	0.70	9.92	7
48	854678	AA630064	40.0	21.23	3.52	1.40	121	2.50	5.93	8
19	362059	AA001431	1.47	5.16	3.51	1.66	5.87	0.68	5.86	8
20	454822	AA677388	0.60	2.09	3.47	0.37	6.64	0.26	29.9	7
23	308696	N92842	0.05	0.16	3.28	1.59	6.55	0.00	4.97	8
22	415788	W84751	1.20	3.83	3.19	0.75	5.82	0.36	5.81	Сĺ
23	489535	AA099262	3.91	12.41	-3.17	5.95	0.84	1,37	4.54	7
77	364865	AA028171	0.80	2.53	3.17	3.39	2.18	4.93	2.18	61
22	740941	AA476288	53.78	169.35	3.15	1.55	5.32	0.41	5.31	n
56	241705	H91680	1.71	5.19	3.04	1.25	4.71	5.51	0.69	7
27	754559	AA406301	1.32	3.98	3.00	2.54	3.88	4.35	1.25	2
28	487327	AA045524	0.49	1.48	2.96	1.51	4.72	0.91	4.72	7
53	643098	AA488676	5.29	15.54	2.94	1.27	2.68	3.2	4.49	63

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64 6	<b>y</b> (	7 (	7	8	7	7	2	7	7	7	7	8	~	7	7	7	0	2	7	~	7	7	7	2	7	70	7	7	7	-	-	<b></b>	-	-		
4.79	4.42	4.95	5.26	4.59	0.00	3.66	3.81	2.47	3.92	4.19	3.45	3.21	3.86	0.87	124	3.83	3.59	4.17	3.58	3.02	3.39	3,49	3,19	3.32	3.63	3.16	ь. 4	3.20	3,19	1.47	1.63	15.38	0.82	0.31		
1.18	4.07	4.85	0.29	0.83	2.12	2.04	1.05	3.41	3.94	96:0	3.06	1.85	3.21	3.08	0.91	0.53	3.21	0.41	1.65	1.72	0.34	0.72	0.97	1.47	3.21	0.53	0.36	0.95	0.51	31.70	4.1	2.08	9.64	0.30		
4.79	1.80	0.86	5.28	4.59	3.14	3.67	3.82	1.28	1.43	4.20	1.25	322	1.22	4.30	3.72	3.86	1.07	4.18	3.59	3.03	<b>7</b> .	3.49	3.16	3.33	0.69	3.16	3.45	3.23	3.20	0.85	1.63	0.71	1.02	17.43		•
86° O	1.32	583	) (2	43 6	 	 		27.5	3.20	0.32	9 9	67.	<u>}</u> :	- 5	S c	) r	2 9	<u> </u>	6.5	0.42	1.40	5.6	, de	030	68.0	138	3	6.0	5 5	) d	0.95 78 at	10.01	0.35 44	0.17		
204	, 60	2.90	787	2.86	2.79	2.6/	2.64	2.50	2.59	2.56	2.50	2.43	2.43	2.35	2.35	2.34	2.34	2.34	2.34	2.31	2.29	2.2	2.16	2.15	51.3	2.10	00.7	36.	1.88	1.86	8.67	5.39	4.69	4.65	4.63	2
•	0.65	4.08	4.26	3.16	1.86	0.02	43.42	1.70	3.12	3.00	19.45	15.91	0.19	8.05	7.96	69.88	1.28	5.70	4.88	48.48	4.15	0.13	1.05	0.44	4.37	4.63	0.92	1.35	0.09	7.71	6.43	3.02	7.68	17.08	7.06	
	0.22	1.41	1.48	1.10	0.67	0.01	16.45	0.65	1.20	1.17	7.79	<b>5</b> .5	0.08	3.43	3.39	29.86	0.55	2.44	2.09	20.98	1.81	90.0	0.49	0.21	2.05	2.20	0.45	0.70	0.03	4.15	0.74	0.56	<u>2</u> .	3.68	1.53	
	N73958	AA456886	AA432049	W46900	AA682386	AA435317	AA428415	W84718	AA701519	AA489743	AA485508	AA411384	R56898	AA598795	AA916325	AA011215	AA010221	AA041499	W42812	AA434502	W20458	R53558	R63646	H24350	H11063	AA452278	AA461174	H86554	AA053684	AA401441	AA455988	N90051	N51498	AA460511	H25546	
	298303	815542	782537	324437	461759	754393	770983	415692	435820	823696	811000	754600	41214	898096	1473304	359835	430231	376518	323238	770880	303099	38959	138788	52071	47418	787938	796806	223350	510136	741977	812074	305809	282104	795809	161456	
	30	ᇊ	32	8	8	. K	3 %	37	38	ස	8	4	42	63	4	5	<b>4</b>	47	8	67	යි	25	22	ß	22	8	æ	23	8	29	8	2	8	B	B	

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

-	-	-	-	<b>-</b>	-	-	_	-	•	-	•-	-		-	-	<b>-</b> -	-	<b>-</b>	-	-	-	~	₩-	-	-	-	-	-	-	-	-	-	-	-	
12.08	0.42	1.00	14.50	13,33	13.47	0.62	1.33	11.93	1.09	7.48	13.64	0.61	2.41	8.86	0.48	0.84	0.20	9.83	0.73	8.89	1.16	0.79	0.45	6. <del>.</del>	1.20	1.27	0.51	6.89	1.71	1.55	1.10	2.15	2.30	2.18	
1,	0.97	13.88	0.84	1.26	. 0.83	12.60	2.26	0.80	12.18	0.80	0.40	1.26	1.81	2.20	0.70	0.66	0.44	0.86	0.34	0.96	0.78	2.68	1.13	<b>1</b> .	1.08	1.00	7.98	2.38	1.72	0.77	2.52	0.62	1.70	1.42	
-	0.92	0.70	1.16	96.0	0.90	0.52	10.78	1,21	1.09	1.48	0.31	11.31	1.97	1.14	12.18	0.92	0.96	0.92	0.73	200	9.88	1.70	1,47	1.30	1.20	1.81	0.57	0.90	7.08	2.28	2.89	2.16	2.01	1,41	
20.5	15.34	1.91	0.51	1.38	0.67	2.12	0.84	1.08	0.62	. 11.18	0.43	1.03	7.91	1.73	0.40	10.97	11.67	<del>2</del> .	11.16	0.87	0.83	6.86	8.81	7.61	8.34	72.7	2.20	0.92	0.56	6.45	4.47	5.87	4.79	5.78	
4 45	4.4	4.37	4.25	4.23	3.97	3.97	3.80	3.78	3.74	3.73	3.69	3.55	3.53	3.48	3.44	3.35	3.32	3.31	3.24	3.18	3.16	3.01	2.97	2.96	2.96	2.84	2.81	2.77	2.77	2.76	2.75	2.70	2.70	2.70	~
Ç	980	3.58	2.10	4.89	4.59	6.59	2.51	3.06	3,20	5.85	18.28	0.32	73.02	3.01	0.21	0.73	0.15	2:94	8,46	0.85	£.5	48.46	6.88	13.27	2.40	0.93	2.99	3.09	4.88	90.0	6.25	32.44	2.47	2.89	
ç	0.09	0.82	0.49	1.16	1.18	1.68	99:0	0.81	0.86	1.57	4.95	0.09	20.71	0.87	90.0	220	0.0	0.89	2.61	0.27	0.48	16.11	2.32	4.48	0.81	0.33	£.08	1.11	1.76	. 0.03	2.28	12.01	0.91	1.07	
A A 701200	AA863204	R63823	AA704718	W69960·	H41222	H19128	AA628410	N48261	AA425382	W38679	AA451886	H06093	N64840	T77410	W47487	N33041	T99719	AA456413	W48762	H38425	H12981	AA136808	AA878880	<b>191197</b>	N47738	H17882	H44861	W31255	R00480	AA160606	AA133721	AA486281	AA598659	AA024898	
67770	1455740	141338	450948	343919	192258	50930	1032774	282117	773157	320857	786657	44351	284701	23790	324313	270343	122782	788386	325128	192401	43207	491001	1493160	307174	277423	50182	176904	306540	123331	592707	512118	842848	897901	365227	
¥	8 8	29	68	69	2	7	22	73	74	22	92	22	82	29	8	<del>8</del>	82	83	R	85	88	87	88	88	8	2	35	ន	8	88	8	26	86	8	

Table 24; Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

_	-	•-	•	-	-	-	•	•	•	•	<b>.</b>		•	•	,	•	·	·	·	•	•	•	•	Ĭ	·	•	•	•-	•	•	,-	-	Ť	•-	
6.67	0.71	1.74	6.57	1.96	0.94	1.75	1,41	75.	2.93	2.01	1.15	2.36	1.80	6.80	3.38	2.70	0.26	0.65	181	133	66.1	07.1	3.50	0.72	0.69	1.24	6.9	1,59	6.65	2.40	. 1.29	0.89	1.26	0.54	
96.0	1.12	1.48	80.0	200	0 60	2.02	80	1.45	2	1.56	12.0	1 72	0.69	. <del>.</del> .		2 64	3	S .	20.1	4, 4 0, 4	- e	- 66	£. c	5.4 5.5	2.5	9	2 2	÷ 5	8 8	8 8	4 92	583	0.53	1.25	
1.47		5.1.5	5 7	4.78	97.1	S. F.	97,۲	67.7	5.45 6.45 6.45	21.1	7 7	÷ ;	£.53	1.0	0.95	16.7	3.5	98.0	1.47	<b>1</b> .90	-28	2.05	3,76	 88.	6.02	25.1	76.7	3. 7	1.1	61.1	0.32 4 38	A7:	. O	4.41	<u>;</u>
+ 57	) i	67:7	6.78	1.92	4.83	6.99	4.00	5.46	4.68	1.46	5.73	6.60	3.52	5.66	0.78	1.13	1.38	8.20	6.74	1.20	5.69	2.68	2.28	3.35	0.80	6.32	3.80	0.78	5.13	0.61	4.58	1.88	96:	8. 0 8. 0	0.00
•	2.69	2.68	2.67	2.66	2.60	2.60	2.56	2.55	2.54	2.54	2.62	2.50	2.50	2.49	2.48	2.47	2.45	2.45	2.45	2.44	2.44	2.43	2.43	2.41	2.39	2.39	2.37	2.37	2.36	2.35	2.34	2.34	2.34	2.33	2.32
	2.38	2.19	2.51	3.75	21.09	0.73	4.25	7.65	3.12	7.59	14.37	13.42	0.07	¥.17	1.57	2.60	5.92	23.38	6.48	7.65	0.13	2.62	3.86	6.17	1.65	4.56	4.66	2.42	0.38	1,74	30.32	26.12	0.54	0. 13	6.68
	0.88	0.82	96.0	7	8.12	0.28	39.1	3.0	1.23	2.99	5.71	5.37	0.03	13.72	0.63	1.05	2.41	9.53	2.65	3,13	0.05	1.08	1.63	2.56	0.69	1.91	1.96	1.02	0.16	0.74	12.93	11.15	0.23	90.0	2.88
	AA701931	AA885311	H29256	N51367	N79738	H75860	AA682631	R17962	W76603	R92601		AA010932	AA676470	AA443004	R15409	T95274	AA865464	AA621031	AA629991	AA434362	AA704844	AA278408	AA778098	AA421389	AA999953	N27992	N76101	AA628146	R12183	AA757732	R98064	R11587	AA621324	W02342	NZ6163
	435567	1461664	49630	283208	289599	230560	431298	32381	345670	196257	361363	359781	882511	811766	53162	120162	1470048	1056203	884514	837908	453218	703930	379920	739237	1636495	265102	299412	1055737	25590	396111	206838	130046	1048794	292071	269411
	100	101	102	8	\$	105	106	107	108	60	110	111	112	113	114	115	15	117	178	5 0	120	5 5	5	123	124	125	126	127	128	129	130	131	132	133	134

-	-	-	-	-	<del>-</del> ,	-	-	<del>-</del>	-	-	-	-	-	-	-	-	-	•	<b>-</b> -	<b>-</b>	-	-	<del>-</del>	-	-	-	-	-	•	-	•	-	-	-
1.28	0.83	0.78	2.08	0.52	0.76	0.43	1.85	0.76	1.46	1.35	3.42	1.52	0.51	0.80	1.86	1.66	0.81	1.55	0.53	3.05	5.49	1.70	5.74	1.18	1.09	1,09	4.10	224	1.13	0.95	2.18	0.92	3.17	0.97
2.50	1.79	1.52	2.27	1.21	5.15	1.55	3.12	1.45	2.79	1.20	1.63	2.50	0.95	0.84	1.71	3.14	1.03	1.44	4.	2.50	1.15	1.33	1.04	1.34	1.52	2.08	1.37	0.92	0.92	0.90	1.84	5.68	1.86	5.76
0.87	2,68	1.74	3.23	1.62	1.23	2.77	1.86	0.75	3.29	1.80	2.82	1.38	1.13	6.45	1.57	1.13	1.02	1.67	1.92	2.59	0.86	0.67	1.13	1.55	2.34	2.19	1.13	224	1.13	1.29	1.52	1.29	1.19	0.97
4.59	3.93	5.17	1.59	5.80	2.02	4.39	2.29	6.15	1.56	4.74	1.21	3.67	6.43	0.88	3.81	2.99	. 6.07	4.18	4.93	0.68	1.28	5.06	0.80	4.62	3.71	3.29	204	3.19	5.41	5.44	3.03	0.67	2.51	0.82
2.34	2.30	2.30	2.29	2.29	2.29	2.29	2.28	2.28	2.28	2.27	2.27	. 2.26	2.28	2.24	2.24	2.23	2.23	2.21	2.21	2.20	2.20	2.19	2.18	2.17	2.17	2.16	2.16	2.15	2.14	2.14	2.14	2.14	2.13	2.13
0	23.45	1.74	8.94	0.13	4.17	19.01	1,33	0.75	4.37	4.23	1,08	960	1,52	76.0	2.12	3.14	97.0	0.05	10.94	3,38	<u>5</u> .	7.68	900	7.96	8.	28.92	5.24	8.31	60.65	1.22	5,69	0.58	11.18	1.14
7 30	10.20	0.76	3.80	90.0	1.82	8.32	0.58	0.33	1.92	1.86	0.48	0.43	0.68	0.42	0.95	1.41	45.0	0.02	4.95	1.53	0.65	3.50	0.02	3.66	0.83	13.37	2.43	3.87	28.28	0.57	2.66	0.27	5.24	0.54
A A 708304	AA197344	AA873089	W76331	AA444111	AA876899	AA011593	H15250	R44776	AA733090	N45398	AA406059	N95435	H25689	N55430	AA598659	AA677295	AA621457	AA047618	AA971641	R83896	AA279409	AA142913	A021434	AA004973	N69876	AA171899	AA134862	AA454165	AA011095	AA419612	H53141	AA777417	AA251386	AA670429
302405	627688	1325751	345055	757135	455179	429678	49687	33096	399577	260187	743026	309929	162077	245936	897901	454798	1055217	376839	1584505	186623	704227	504420	364173	428928	297682	610374	502436	795322	359661	746148	202621	449061	684562	878836
125	3 8	137	138	139	40	141	142	143	<u> </u>	145	146	147	48	149	150	151	152	153	<del>2</del> 2	155	156	157	158	159	160	161	162	<u> </u>	<del>2</del>	165	166	167	168	169

Table 2.4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

<del>-</del> .	<b>-</b>	•	<b>-</b>	-	-	٠.	- ,	_	-	-	-	-	•	•		- •	- ,	-	-	-	_	-	-	-	-			- •	- ,	<b>-</b> ,	-	-	-		-	-	_	•	
4.99	4.82	0.84 48.0	1.32	1.89	2	3 7		3.11	2.79	1.39	0.25	3.92	17	8 F.	2 8	79.0	) 5. G	0,15	1,50	1.36	0.84	1.12	1.74	<u>5</u>	2.64	107	, ;	, v.	/٥٠٢	0.59	0.69	2.01	0.65	. 0.65	0.68	1,39	0.62	3	
1.09	1.69	1.07	4.86	000	9 6	1.65	1.47	1.24	3.06	1.24	2.62	121	9	9 6	3.50	1.09	0.71	0.13	1.68	3.03	0.70	2.17	1.67	157	5 6	21.0	3.10	0.99	1.23	<u>.</u>	1.49	66:0	1.52	1.72	1.53	4 20		15.1	
1.33	0.88	1.29	780	; ;	80:	1.10	<del>2</del> .	1.65	1.16	36.1.	i K	3 6	9 1	0.73	1.73	1.05	0.60	7.72	1.29	1.36	5.44	1.13	7		3	5.2 2.	2.82	2.51	4.23	1.63	1.81	2.01	1.97	3.14	2.08	÷	<u>-</u> -	7	
1.07	1.07	6.25	160	8:	3.70	0.89	4.19	2.34	1.31	2.7		£. :4	1.94	5.14	1.18	5.47	0.93	0.20	3.72	2.42	÷ +	2 .	7 7	3.01	48.0	0.16	1.27	4.18	0.98	4.55	4.11	3.08	3.94	257	3.70	, c	1.28	4.88	
2.12	. 212	; ;		<b>1.3</b>	2.10	2.10	2.09	2.08	a c	) i	2.07	2.07	2.07	2.07	2.06	2.06	2.05	2.05	2.05	75.	7 7	<b>7</b> ,0 <del>4</del>	2.04	2.04	2.03	2.03	2.83	2.03	2.03	2.03	2.02	202	20 2	1 8	20.6	2.02	2.02	2.01	w.
0.78	2	#0.83	5.23	13.63	13.41	3.21	2.34	27.88	32.5	2.90	0.79	36.21	69.0	1.92	3.48	2.14	16.88	961	5 2	2 5	50. 10.	0.27	2.06	3.34	0.75	18.02	13.69	4.76	0.77	5.26	4.26	4. A	3. 4. 2. 4.	70:07	) i	7.47	3.17	0.24	
75.0	5	42.83	2	6.45	6.39	1.53	1.12	42.37		97.	0,38	17.47	0.33	0.93	1,69	2	8.23	4 60	200	<b>1</b>	20.0	0.13	<u>5</u>	<del>7</del> 9.	0.37	8.88	6,75	2.35	0.38	260	2.11				<b>5</b>	3.71	1.57	0,12	
2302044	AA/ 06833	W95187	H83283	AA626242	AA010223	AA521414	NAZZOB	AAAEEEE	AA455363	AA453040	AA676948	AA876375	N71769	AA187789	AA030046	W58013	R45964	. A A D 3 4 E 4 3	AA031313	AA011183	AA430242	H09996	T61792	AA676840	AA872602	AA151002	AA968896	N20328	AA598572	AAROOR3	AA164782	A 4 5 0 5 0 7	A 19050/	00/0444	H67292	AA490985	R10948	H09749	
000	451888	361668	199175	745524	430235	826099	307608		813438	788574	460189	1493107	290702	624379.	469969	341336	35575	7000	4/0393	21/805	781091	46916	78946	460114	1323328	504927	1674594	264597	897822	4044078	505001		20/104	324563	229573	824588	129148	46453	
,	2 ;	17	172	173	174	175	140	2 [	) <u> </u>	178	179	<u>8</u>	181	182	183	18	, 2 , 2 , 3	3 5	9 9	<u>\8</u>	188	189	190	191	192	193	4	195	40,	5 5	100	2 0	660	3	201	202	. 203	204	

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

-	-	-	-		-	-	_	-	-	-	-	-	-	-	-	-	-	<del>-</del>	-	-	-	-	<del>-</del> -	-	-	-	-	-	-	-	-	-	-	-	
¥	1.24	<del>1.</del>	0.80	0.79	0.59	3.62	1.46	1.11	0.80	3.20	1.30	0.75	0.51	1.50	1.78	1.31	0.73	<del>1</del> .30	6.57	2.08	0.82	1.01	0.97	0.93	3.07	1.09	68'0	1.10	0.93	1.65	1.13	0.97	2.39	0.09	
1.35	2.50	1.20	1.69	1.29	1.17	1.20	4.84	3.49	1.22	1.25	1.07	1.82	1.38	1.09	1.28	3.99	2.79	3.96	0.42	2.03	1.05	0.93	1.03	1.19	2.59	1.15	0.76	1.00	1.26	3.06	4.27	1.19	3.27	0.67	
1.73	0.76	0.98	1.88	4.31	5.08	1.38	0.80	2.01	1.69	<u>.</u> .	1.65	0.76	0.95	1.33	1.5	1.33	3.06	1.63	0,40	364	3.99	1.02	0.98	9.	0.82	1.10	2.06	200	1.74	1.80	1.14	1.77	1.18	6.77	
360	3.49	4.80	3.49	1.56	1.09	1.71	0.81	1.29	4.10	2.02	3.98	4.53	5.03	3.92	3.22	1.40	1.22	0.87	0.36	00.0	1.86	4.75	4.71	4.62	1.20	4.33	3.95	3.55	3.70	1.10	1.07	3.66	0.75	0.05	
00 6	700	1.99	1.99	1.99	1.98	1.98	1.98	1.98	1.98	1.98	1.97	1.97	1.97	1.96	1.96	1.95	1.95	1.94	1.94	1.93	1.93	1.93	1.92	1.92	1.92	1.92	1.91	1.91	1.91	1.90	1.90	1.90	1.90	1.89	1
7.0 7	9.35	0.91	19.17	2.63	0.16	10,70	3.81	4.90	8.22	3.66	0.25	0.07	0.09	2.75	1.71	1.89	33.80	1.30	1.51	0.02	4.71	0.54	1.01	0.41	4.14	99'0	1.96	55.28	6.25	1.81	0.56	8.94	3.29	4.03	
, 03	4.68	0.45	9.63	1.32	90.0	5.41	1.93	2.48	4.10	1,85	0.13	9.0	9.	04.	0.87	0.97	47.34	0.67	0.78	0.01	2.44	0.28	0.53	0.21	2.16	0.35	8.	28.91	3.28	0.95	0.29	4.71	1.73	2.13	
703667	N51382	R31114	AA195601	R07948	W02073	N26008	AA879435	AA664105	W15319	N62293	AA455825		W32729	AA259115	AA169173	H38263	AA399074	AA399433	AA434454	H04828	AA634379	T7007TT	H29290	R39364	AA449048	H20717	R01245	AA453105	AA478606	H03146	W79069	R92801	R23609	171061	
70000	283191	134265	665127	127063	327247	268837	1505688	855634	322643	290346	811609	69272	327495	686664	610006	190950	729953	726699	770319	43966	743828	23219	49858	24632	785816	51420	124137	789091	753663	151597	346966	197056	132140	84078	
300	300	207	208	209	210	211	212	213	214	212	216	217	218	219	220	221	222	223	224	225	226	227	228	229	230	231	232	233	234	235	236	237	238	239	

39**8/852** 

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3.84 4	1.20	3.33	0.61	0.86	1.70	1.01	121	0.89	3.29	1.22	1.05	2 60	5.28	3.29	1.57	190	3.13	45.	900	990	4 47	9.6	5 6	0.63	4	55.	0.71	1.30	1.21	4.80	1.39	0.92	3.23	3.61	<b>!</b>	
1.85	1.19	1.50	1.46	4.52	4.05	138	7	2.50	2.5	<u> </u>	5	\$ 4 4	2 4	S &	7.02	3.09 1.84	<u> </u>	7.93	7 ;	: ;	27.1	<u>.</u>	S 6	40.0 40.4	90 +	20.1	<u>†</u> <del>1</del>	3.76	3.38	06:0	3.02	4 79	77.		<b>S</b> 0.2	
1,03	1.68	1.70	3.16	60.	2 0	6.50	 	0.1	0.97	1.5.T	55.	3.06	9.4	0.73	1.46	0.71	1.62	\$ 10	1.65	1.46	1.44	1.22	1.28	5.52	1.07	3 5 C	193	1.22	1,77	180		5 C	0.32	0.92	07:1	
0.85	3.48	-	2 6	9.5	<b>s</b> .	1.20	3.06	3.38	1.83	1.07	9.83	3.78	0.26	0.46	0.93	1.17	3.18	1.30	3.23	3.64	3.35	3.26	1.53	5.61	4.15	3.92	3.22	200	8 5	n 4	0.0	1.26	3.41	9.40	1.02	
1.89	68	6 6 6 6	00.	88.	<del>.</del>	1.88	1.85	1.85	1.85	1.84	1.84	1.83	1.83	1.83	1.83	1.82	1.82	1.82	1.81	1.79	1.79	1.79	1.78	1.78	1.78	1.78	<u>r.</u>	7.7	);;	1.77.	1.77	1.76	1.78	1.75	1.74	∞
2 + 5		84.7	0.89	53.13	2.13	2.63	3.49	8.28	1.09	1.38	41.96	0.53	5.23	1.90	1.56	2.99	2.48	2.68	12.19	2.78	2.18	2.31	2.53	0.03	2.80	8,35	0.87	4.09	7.89	2.64	5.15	0.26	7.01	2.78	10.90	
ţ	7 .	3.97	0.47	28.27	1.4	1.40	1.88	4.47	0.59	0.75	22.82	0.29	2.86	<u>5</u>	0.86	1.64	1.36	1.48	6.74	.55	1.22	1.29	1.42	0.0	1.57	4.81	0.49	2.3	4.46	1.49	2.82	0.14	3.99	1.59	6.25	
1070	16161	AA453531	AA682321	AA443698	AA453863	R98344	AA155787	AA476961	N20989	AA704802	AA664406	AA609982	R38645	AA772497	AA677984	191744	AA282594	AA434117	AA677397	W88623	AA620421	AA448194	N46759	AA885052	R71124	AA030004	H61082	AA620611	R53342	AA905838	AA827405	AA126662	R70684	AA876354	W05033	
	78680	788136	462926	784016	813719	201641	590253	739578	265853	452466	868652	1031203	23000	413300	430763	283328	713019	770216	454896	417691	951022	782797	277112	1466485	142944	470218	206172	1048702	40031	1505294	1422794	502593	141815	1493079	298648	
•	240	74	242	243	44	245	246	247	248	249	250	251	252	253	25.	255	256	257	25.5	250	260	284	262	263	264	265	266	267	268	569	270	271	272	273	274	1

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0.89	1.38	1.28	0.48	1.07	3.13	0.81	0.28	3.32	121	0.40	0.71	3.71	0.73	4.33	3.77	3.29	. 0.53	3.62	9.0 48.	3.67	0.83	1.05	0.25	0.45	1.26	3.00	1.76	0.16	3.43	1.35	-1.17	0.72	0.99	1.24	
1.11	0.81	0.93	1.10	1.53	1.38	1.81	0.87	1.21	3.37	1.42	1.42	1.47	5.1	1.21	1.07	141	4.89	1.33	1.09	1.42	0.95	0.99	0.76	1.78	1.12	1.46	0.62	0.91	1.33	1.31	1,15	1.65	1.23	1.01	
1.24	1.36	1.28	2.26	0.59	1.60	3.25	2.31	1.20	1,22	1.45	3.33	0.78	1.43	0.78	1.18	0.65	0.51	0.95	1.63	1.03	0.57	1.50	4.85	3.15	1.26	£.	0.87	2.51	1.01	0.81	1.18	121	0.80	1.11	•
3.73	3.43	3.48	3.11	3.75	0.81	1.05	3.45	1.16	1.08	3.62	1.41	0.91	3.01	0.53	0.83	1.50	0.91	0.94	3.27	0.70	4.47	3.29	0.95	1.42	3.17	1.22	3.52	3.15	0.95	3.23	3.18	3.11	3.66	3.31	
1.74	1.74	1.74	1.74	1.74	1.73	1.73	1.73	1.72	1.72	1.72	1.72	1.72	1.72	1,71	1.71	1.71	1.7.	1.71	1.71	1.71	1.70	1.70	1.70	1.70	1.70	1.70	69.1	89.1	1.68	1.68	1.67	1.67	1.67	1.67	
2.27	5.31	4.53	3.02	29.04	2.13	57.00	5.19	1.38	0.58	35.33	5.05	1.42	7.12	5.74	0.69	2.60	0.82	1.74	62.95	2.09	0.33	21.50	5.98	3.24	1.78	3.16	90.0	0.07	0.90	10.94	47.59	5.91	8.82	0.85	
1.30	9. 20.	2.60	1.74	18.74	1.23	32.95	3.00	0.80	9.34	20.52	2.94	0.82	4.15	3.35	0.40	1.52	0.48	1.01	36.82	1.23	0.19	12.62	3.51	1.91	1.05	1.86	0.03	9.	0.54	6.53	28.46	3.54	5.28	0.51	
AA417344	T81289	. AA457235	AA057032	R09179	AA455980	. AA620995	R38943	W71990	N26608	AA459761	AA255551	H80063	W73724	R81831	AA700843	AA701948		AA464598	AA047260	W86124	R54179	AA456695	A1003028	H96392	H94563	R82317	H84734	H73806	AA437093	AA464068	AA664389	AA733061	AA420992	AA883114	
731168	25061	838668	488913	127610	812053	1056172	24958	345701	269172	796366	682477	249070	344108	147834	436097	435919	25122	812961	488964	416227	41608	813149	1684274	256680	243159	149058	220135	214713	757325	810284	868630	389562	731255	1466942	
275	276	277	278	279	280	281	282	283	284	282	286	287	288	289	290	284	292	293	294	295	296	297	298	299	300	30	302	303	304	305	306	307	308	309	

Table 2-4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

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3.30	1.22	0.71	3.88	1.61	1.38	1.38	0.87	17	0.62	3.05	1.68	0.34	990	1.58	78.0	5.54	78	1 48	. c		- 6	6.0	6.60	2 E	8 6	- a		02.0	2,6	1 24	2 0	n 4	0.55	1.3¢	9.1	
1.25	108	1.09	1.28	£ 5	£ 5	3 5		, t	£ .	36.1	<u> </u>	Į Ę	2, C	4.24	<u> </u>	3 6	200	79.7	B 6	0.08	52	4.10	1.23	9. 7.	18:0	35.5	9.68	2 .	26.	<u> </u>	0.37	88.0	0.41	15. 1	(S.9)	
0 03	125	į (	9 e	9.0	5 6	0.98	5.6	3.00	07.1	1.40	1.5/	0.58	2.18		6.0	<b>\$ \$</b>	6.0	1.20	3.28	1.15	0.98	0.77	1.45	<del>2</del> .	0.88	25 E		C7.1	3.05	1.38	1.24	0.81	4.59	0.58	0.88 8	
	0.46	0.00	3.80	0.80	0,00	3.30	8 !	1,27	3.21	3.23	0.78	3.06	3.22	3.72	3.08	3.80	60.0	3.52	0.80	3.88	3.11	0.71	3.08	3.61	0.73	2.	3.39	3.03	0.99	3.12	3.26	3.83	0.74	3.04	3.24	
1	1.67	1.67	1.66	1.66	1.86	1.65	1.65	49. 49.	1.64	1.64	1.64	1.63	1.63	1.63	1.62	1.62	1.62	1.62	1.61	1.61	1.61	1.61	1.60	1.60	1.59	1.59	1.59	1.59	1.58	1.58	1.58	1.58	1.57	1.56	1.55	10
	0.23	0.67	0.15	42.64	2.46	2.82	0.59	4.02	0,80	7.92	0.71	0.28	12.12	0.76	9.47	5.76	12.39	17.82	0.89	3.93	3.25	2.10	4.80	0,33	5.82	32	2.85	26.88	227	07.7	2.42	11.90	1.36	1.75	1.78	
	0.14	0.34	0.09	25.66	1.48	1.70	0.35	2.44	0.49	£83	0.43	0.17	7.43	0.46	5.83	3.54	7.64	11.02	0.55	2.45	2.02	131	3.00	0.21	3.72	2.03	1.86	16.93	<del>5</del> .	4.86	<del>2</del> .	7,56	98.0	1.12	1.14	
	AA884397	AA788738	AA983362	AA757417	<b>T65398</b>	R89358	H48138	H16439	AA719257	AA479512	AA279990	H15040	AA504239	H58655	R43553	AA150422	R43053	AA012867	AA459472	AA774665	W73748	AA487676	AA070435	N48913	AA156926	AA406019	AA453900	AA116061	R37357	AA169606	H11036	AA279145	AA458377	H24580	H23548	
	1461725	1240347	1592715	395417	21634	196038	183780	47527	1292593	754313	705064	48454	825031	212325	32794	491415	32509	360885	814460	856289	344272	841624	530954	279613	502518	742132	813755	531862	28270	594266	47378	703943	813256	160730	52002	
	310	311	312	313	314	315	316	317	318	319	320	321	322	323	324	325	326	327	328	3.0	330	3 8	330	333	3 5	335	336	337	338	339	340	7	<b>3</b>	; <del>S</del>	***	

Table 2-4; Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one turnor sample.

345	204251	H59188	1.38	2.13	55. 58.	3.28	1.16	0.93	3.45	
347	712888	AA282159	6.43	6.93	<u>.</u>	3.59	1.45	. 99'0	0.47	_
848	951108	AA620466	204	3.15	<u> </u>	0.58	0.79	0.86	3.94	•
349	743187	AA400002	1.45	2.24	7.	1.00	0.70	0.68	3.79	_
350	66952	T69522	0.1	0.16	7. 2.	0.99	0.71	3.21	1.26	_
351	129589	R16541	0.92	7.	1.54	0.59	3.16	1.98	0.42	_
352	461699	AA682274	5.46	8.37	53.1	3.63	8.	0.75	0.72	_
353	452676	AA779225	0.40	0.61	1.52	0.71	1.06	3.08	. 178	•
354	196992	R93124	1,4	2.19	1.52	0.64	3.67	0.94	0.83	•
355	1501546	AA886792	9.0	1.42	1.52	0.86	3.03	1.48	0.70	_
356	754591	AA406320	3.70	5.60	1.51	0.83	0.54	0.56	4.14	_
357	703827	AA278842	1,39	5.09	1.51	0.78	<b>6</b> .	<del>1</del> .13	3.11	_
358	1486752	AA912071	1.05	1.58	1.50	0.69	0.83	0.71	3.78	_
359	281045	N50904	0.98	1.47	55.	0.93	0.85	3.38	0.85	•
360	505414	AA161248	90.0	0.0	1.47	0.78	3.03	1.63	0.43	_
361	179631	H51122	0.50	0.73	1,47	0.68	1.09	0.74	3.36	•
362	201125	R98472	9.68	14,17	1.46	3.64	1.24	92.0	0.22	•
363	753271	AA411668	<u>5</u>	1.51	1.46	0.94	0.87	0.95	3.08	_
36.	46844	H10073	0.11	0.16	1.45	3.45	0.79	0.80	0.78	~
365	453988	AA779212	0.48	0.70	1.45	3.31	8.	0.67	0.80	_
366	141548	R73088	1.37	1.98	1.44	0.86	0.70	0.51	3.69	Ψ-
367	151184	H02294	2.40	3.43	1.43	0.35	0.85	0.77	3.76	-
368	263697	H99672	8.21	11.76	1.43	3.35	1.55	0.52	0.31	~
369	845521	AA644563	<u>5</u> .	2.63	1.43	3.26	0.74	1,12	0.61	<b>-</b>
370	428796	AA004671	90.0	0.0	1.43			0.91	0.86	_
371	124567	R01937	90.0	0.11	1.43	3.11	92.0	4.14	0.69	•
. 372	44300	H06377	2.96	4.22	1.42	1.02	0.50	9:0	3.22	~
373	454564	AA677025	0.87	1.23	1,41	3.29	0.79	0.81	0.76	~
374	196612	R92994	0.45	0.63	1.40	3.60	0.87	0.59	0.56	-
375	665774	AA193254	11.47	16.09	1.40	3.04	1.19	0.60	0.78	~
376	416305	W86183	0.80	÷	64.1	0.88	3.65	0.57	0.50	~
377	773483	AA428014	9.13	12.76	1.40	3.08	1.03	0.71	0.77	-
378	244310	N54788	0.57	0.80	<del>4</del> .	3.47	1.05	0.48	0.58	<b>-</b>
379	195751	R89082	1.31	1.83	1.39	3.29	1.17	0.74	0.37	Ψ-
						Ξ				

Table 2.4: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or higher in any one tumor sample.

-	-	-	-	٠,	_	<b>-</b> -	<b>-</b> -	<b>-</b>	-	-	-	-	-	-	-	-	-	-	-	-	-	
0.89	3.23	3.78	100	5	0.78	0.75	0.51	0.39	0.70	78.0	0.49	0.42	0.52	0.67	0.40	0.45	0.37	0.26	09:0	0.40	0.16	
0.82	0.69	540	4 6	2.0	3.28	0.80	0.49	25.0	0.36	07.0	220	) ()	, e	5 6	<b>4</b> .01	0.85	38	95.0	0.00	0.30	9 6	?
3.15	Cac	3 6	0.5/	<del>.</del> Ş.	0.78	3.48	88.	3 i		) t	90.0	9 9	E).1	U.30	97.0	0.45	) ; ;	5,63	0.75	0.23	G .	37.5
0.53	7.	6.7	0.78	3.35	0.53	3 6	, 5 (	0.40	0.66	3.50	6.59	3.08	3.11	0.85 5	3.12	Q.13	3.10	0.47	3.25	3.29	0.12	- 5
38	9	1.36	1.36	1.36	36	<u> </u>	<b>5</b> .	1.33	1.32	1.32	1.30	1.29	1.29	1.25	1,25	1.25	121	1.21	1.20	1.09	1.05	
9	70.75 70.75	<u>.</u> 2	2.34	14 94	<u> </u>	06.4	1.52	3.71	1.41	0.57	1.30	6.27	9.28	2.54	7.36	0.10	6.03	2.20	4.23	25.11	14.06	1
į	2.61	<del>.</del> 38.	1.72	**	70.1	3.20	1.13	2.79	1.07	0,43	1.01	4.08	7.18	2.03	5.89	90.0	4.98	1.82	3.53	23.01	13.33	
	AA922309	AA706969	W93483	2460074	AA588810	173883	N49574	N49629	AA102670	AA040265	AA478542	AA074677	H85806	N53421	R67147	AA443706	H49519	AA401137	AA058597	AA843592	AA705516	
•	1474284	451911	357284		897952	84484	277708	243741	563598	378043	784772	544639	222157	284100	42373	784017	178860	741497	380890	1390584	462325	
	88	381	282	3 6	383	384	385	386	387	388	389	390	391	392	393	38	395	396	397	398	399	

Table 2-5: Poor vs. Good Clinical Outcome: Genes upregulated 3 fold or more in at least 2 tumor samples

9;	st 2	
Tabld 2-5: Poor vs Good Clinical Outcome:	Genes upregulated 3 fold or more in at least 2	tumor samples.

Order	IMAGE Clone ID	Accession No.	AveGood	AvePoor	Fold(AvePoor/ AveGood)	Fold(AvePoor/ Fold(mpm31/Av Fold(mpm32/ AveGood) eGood) AveGood)		Fold(mpm33/A	Fold(mpm33/A Fold(mpm34/Av # of Tumor veGood) Samples	# of Tumor Samples
•			,	4, 4,	20 71	2.52	57,56	34.98	23.80	ы
<del>,</del>	753078	AA436478	7.42	5.10		690	16.64	3,34	3.64	6
7	124071	R02586	3.5	3.21	9 1	896	7.75	1.51	27.7	n
m	346321	W79647	9.5 49. ;	3.29	9.1/	3.08	6.82	0.22	6.80	m
4	855624	AA664101	24.18	108.55	84.4 10.4	. 4 4 80	5.91	4.66	0.57	•
S	42325	R61073	13.03	27.76		S 6	3.29	1.30	3.29	80
တ	287528		2.13	6,32	/6.7 	5	361	3.01	3.60	•
7	756595	AA444051	 8-	රා භ	2.6/	2 6	5 5	0.37	62.14	7
œ	452374	AA700876	9.0	19.98	31.26	† * * * * * * * * * * * * * * * * * * *	25. AC	12.28	0.86	. 7
တ	357105	W93315	0.87	8.54 42	9.8	10.0	2007 100 100 100 100 100 100 100 100 100	182	0.33	7
9	460798	AA708152	1.24	6.95	5.61	0.47	17.80	3 5	\$ 17	. 2
F	284601	N76201	0.27	1.50	5.49	0.82	10.01 10.01	2.0	# # # # # # # # # # # # # # # # # # #	. ~
12	460487	AA677706	53.51	237.74	4.44	0.02	8.8	3 ;	200	, ,
<u>£</u>	248412	N58558	0.86	3.63	4.25	0.95	7.66	0.75	5 }	• (
4	212188	H68848	0.58	2.37	£.23	0.93	6.76	2.47	6.75	۷ (
. <del>1</del>	782513	AA432030	4.08	16.09	3.86	1.06	1.68	6.19	6.98	7 (
<u> </u>	782513	AA432030	2.03	7.78	3.83	1,37	1.26	6.00	6.70	<b>N</b>
2 5	129128	R10935	0.38	1,47	3,81	0.92	3.71	0.70	9.82	7
- ¤	854678	AA630084	804	21.23	3.52	1.40	4.21	2.50	5.95	79
5 5	262050	AA001431	1.67	5.16	3.51	1.66	5.87	99.0	5.86	7
2 6	454822	AA677388	0.60	2.09	3.47	0.37	6.64	0.26	6.62	7
7 6	308698	N92842	0.05	0.16	3.28	1.59	6.55	0.00	4.97	P) (
3 :	415766	W84751	25	3.83	3.19	0.75	6.82	0.38	5.81	7 (
1 2	489535	AA099262	3.9	12.41	3.17	5.95	0.84	1.37	4.54	7
24	364865	AA028171	0.80	2.53	3.17	3.39	2.18	4.83	2.18 8	N .
, נ נ	740044	AA478298	63.79	169.35	3.15	1.55	5.32	0.41	5.31	7
3 8	344705	Latero	7.	2,5	3.04	1.26	4.71	5.51	0.69	~
9 7	784850	A 4 406301	. 2	3.96	3.00	2.54	3.88	4.35	1.25	2
ž č	487427	AA045524	0.49	1.46	2.96	1.51	4.72	0.91	4.72	7
3 8	94300g	A A 4 8 8 6 7 B	5.29	15.54	2.94	1.27	2.66	3.34	4.49	7
3 8	298303	N73958	0.22	0.65	2.94	0.99	4.79	1.18	4.79	~

64	7	7	~	7	8	2	. 6	~	7	. 2	. 2	. ~			, ,	4 (	<b>y</b> (	7	7	8	7	7	8	2	۰ ۲		7	7	7	~	
4.42	4.95	5.28	4.59	0.00	99	1.81	2.67	3.92	4 49	3.45	3.5	98.	0.83	5 6	\$ 7. C	3.83	3.59	4.17	3.58	3.02	3.39	3.49	3.18	132	20.0	9 9	3.16	3.44	3.20	3.19	
4.07	4.85	0.29	183	3 13	<u>.</u> 5	t 8	8 4	70, 6	, d	<b>5</b> 5 6	5.00		3.21	3.08	0.91	0.53	3.21	0.41	1.65	1.72	7	54.0	70.0		) d	3.21	0.53	0.36	0.95	0.51	•
1.80	0.86	38	2 5		* ! 	3.67	3.82	37.	3.	4.20	1.25	3.22	1.22	4.30	3.72	3.86	1.07	4.18	3.59	203	300	<b>t</b>	0.4	2 6	3.33	0.59	3.16	3,45	3.22	120	<b>.</b>
1.32	0.83	8. 8	0.02	5.7	5.43	1.19	1.72	3.20	0.92	99.0	1.99	1.43	Ξ.	1,13	3.49	1.13	1.48	0.57	0.42	· •	2 (	1.78	<b>4</b> 6.0	1.28	0.39	0.89	1.38	0.44	0.16		CC'O
S	<b>7</b> .90	78.7	2.86	2.79	2.67	2.64	2.60	2.59	2.56	2.50	2.43	2.43	2.35	2.35	2.34	2.34	2.34	75.0	, i	2 6	82.7	2.21	2.16	2.15	2.13	2.10	2.06	1 93	) Q	90:-	1.86
	80.4	4.26	3.16	1.85	0.02	43.42	1.70	3.12	3.00	19.45	15.91	0.19	8.05	7.96	69.88	1.28	5.70		4.00	40.40	4.15	0.13	1.05	0.44	4.37	4.63	0.92	7 7	2 6	<b>3</b>	7.71
;	1.4.	1.48	1.1	0.67	0.01	16.45	0.65	57	1.17	7.79	6.54	90.0	3.43	3.39	29.86	0.55	2,44	‡ 6	50.7 50.7	20.32	 	90.0	0.49	0.21	2.05	220	0.45		2 6	8	4.15
į	AA456886	AA432049	W46900	AA682386	AA436317	AA428415	W84716	AA701519	AA489743	AA485508	AA411384	R56898	AA598795	AA916325	AA011215	AA010221	4404400	AAU41493	W42812	AA434502	W20458	R53558	R63646	H24350	H11083	AA452278	AB451174		H86554	AA053684	AA401441
	815542	782537	324437	461759	754393	770983	415692	435820	823696	811000	754600	41214	898098	1473304	359835	42024	10000	3/65/6	323238	770880	303089	39959	138788	52071	47418	787938	705906	00000	223350	510136	741977
	9	32	33	8	35	ဗ္ဗ	37	88	9	8 8	4	42	43	: 4	7	2 4	<b>?</b>	4	<b>4</b>	49	20	51	25	53	3	ָ ע	3 4	B 1	21	28	23

Table 2-6: Poor vs. Good Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Table 2-6: Poor vs Good Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples

Order	IMAGE Clone Id	Accession No.	Tissue Distribution	AveGood	AvePoor	Fold(AvePooriA Foldimpm31/Av Fold(mpm32/ Fold(mpm34/ # of Tumors veGood) AveGood) AveGood) Samples	Fold(mpm31/Av eGood)	Fold(mpm32/ AveGood)	Fold(mpm33/A veGood)	Fold(mpm34/ AveGood)	# of Tumors Samples
-	753076	AA436479	thymus	1.42	42.16	29.71	2.52	57.56	<b>%</b> .88	23.80	ಉ
7	452374	AA700876	prostate	0.64	19.98	31.26	0.24	62.30	0.37	62.14	. 11
ю	357105	W93315		0.87	8.54	9.81	0.81	25.33	12.26	0.89	~ ~
4	284601	N76201	brain	0.27	1.50	5.49	0.82	10.19	0.77	10,17	8
\$	346321	W74079	spleen	. 79.0	3,29	5.17	3.69	7.75	1.51	7.73	8
9	855624	AA664101	testis	24.18	108.55	4.49	4.12	6.82	0.22	6.80	~~
7	460487	AA677706	bone	53.51	237.74	4.44	0.02	8.87	0.03	8.85	7
80	248412	N58558	livar	98.0	3.63	4.25	0.95	7.66	0.75	7.64	7
თ	212188	H59148	liver	0.56	2.37	4.23	0.93	6.76	. 2.47	8.75	7
5	782513	AA432030	testis	4.08	18.09	3.85	1.06	1.58	6.19	8.98	74
¥	782513	AA432030	testis	2.03	7.78	3.83	1.37	1.26	6.00	6.70	8
12	362059	AA001431	pone .	1.47	5.16	3.51	1.66	5.87	0.66	5.88	7
₽ £	454822	AA677388	colon	0.60	5.09	3.47	0.37	. 6.64	0.26	6.62	7
4	415766	W84751	heart .	1.20	3.83	3.19	0.78	5.82	0.36	5.81	7
15	740941	AA478298	testis	53.79	169.35	3.15	1.66	5.32	0.41	5.31	8
16	324437	W46900	bone	1.10	3.16	2.86	0.62	5.28	0.29	279	7

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

Table 2-7: Good vs Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

		Gen Bank			Endia ve Goodia Foldimpm 27/Av Foldimpm 28/Av Foldimpm 28/Av Foldimpm 30/Av Foldimpm 30/Av	Fold(mpm27/Av F	old(mpm28/Av l	old(mpm28/Av F	old(mpm30/Av Fc	old(mpm36/Av	# of Tumors
Order	Clone	Accession	AveGood	AvePoor	vePoorl	ePoor)	ePoor)	ePoor)	ePoor)	ePoor)	Saidues
		Number					1	i	830	7.30	*
	200679	W05628	2.55	0.50	5.07	7.07	6.80	(v.)	6.63	8.8	4
- (	B44482	AA486471	28.87	8.85	4.21	0.70	3.61	6.33	- C	24 45	· e7
e es	471641	AA034945	11.03	1.08	10.17	21.19	2.20	27	7.10	40.4	) e2
•	040490	N47717	34.87	4.42	873	22.87	<del>5</del> .	0.27	9	2 6	•
<b>;</b>	287089	N48259	27.13	3.45	7.87	22.41	0.41	0.67	2.02	0.00 45 90	<b>9</b> 67
) נו	155072	R71393	3.32	0.45	7,44	8.50	1.97	2.30	70.5	10.50	) e:
o	208210	H65286	2.70	0.42	6.48	14.00	1.98	0.98 0.0	000	20.50	
. α	770056	AA430545	26.72	4.62	5.78	12.05	0.67	1.90	0.00	12 92	, en
<b>,</b> c	747050	AA282273	38:22	6.94	5.50	8.84	25.	0.35		1.2A	, e3
• \$	81331	T60111	4.13	0.82	5.03	11.18	.30	0.80	0.0 40	5.42	· m
÷	287778	N59158	3.49	0.71	6.90	14.39	0.35	0.46	90.0	7.7	•
: 2	41228	R58853	0.05	0.0	4.38	9.07	2.53	4. 0 0 t	1 9	3.18	, m
E	122345	T99191	0.03	0.01	4.16	12.66	4.1/	97.0	900	34	က
2 7	186918	H43317	6.27	1.60	3.83	3.28	1.41	3.	) i	5.03	m
<b>.</b> \$2	1049033	AA778675	2.50	0.65	3.87	4.86	2.38	4, c	2.55 5.55	6.23	· (*)
<b>\$</b>	195385	R88904	18.	0.50	3.62	6.03	1,45	88.0	2.0	25.0	· «1
2 \$	50114	H16743	1.48	0.43	3.43	5.36	4.18	4.79	97.7		· en
= \$	038644	AA456975	87.41	25.99	3.36	5.71	4.22	4.22	25.1	<u>.</u> 5	· "
<u>•</u>	282408	N45282	1. 6	1.55	3.32	7.31	0.87	1.42	3.24	7	) r
e 6	745576	A4626248	418	1.27	3.28	5.85	1.7	0.34 4	0.7	,	. "
3 ;	145067	H15928	5	2	2.82	2.50	3.69	3,70	1.59		? r
<b>1</b>		AA779728	76.12	27.08	2.81	2.22	3.21	3.87	0.79	9.6	) r
3 8	438444	R68272	202	7.0	2.75	4.16	1.66	1.35	3.32	2.20	י רי
3 2	270499	AA777187	5	71.7	2.37	3.23	1.20	66.O	3,13	 	, c
<b>5</b> 2	201010	AA424535	£ 28	0.37	11.65	1.60	25.69	28.44	24.5	<u>,</u>	, ,
3 2	707737	N95228	10.81	98'0	11.29	1,65	7.47	44.45	.65.	4:	• 6
3 5	\$03628	AA134576	17.77	0.80	8.08	1.44	1.16	33.58	3. 6	; ;	
8	41869	R66438	7	0.28	5.03	1.48	5.93 1.93	15.74	) ()	60	1 64
28	301735	N90882	4. 6.	0.80	4.56	8.05 50.	2.75	3 5		2.65	~
8	136805	R35051	32.22	7.39	4.36	2.55	3.53	11.13	- 4 - 4 - 4 - 4 - 4 - 4 - 4 - 4 - 4 - 4	37.	1 7
8	362278	AA001219	5.53	1,41	3.9	2.64	<b>3</b>	9 9	9 6	5	7
3	758708	AA443803	. 2.58	0.67	3.84	0.50	4.59	90.03	6.3	86.0	2
2	285028	N21084	1.22	0.32	3.78	<del>-</del> 8:	<b>4</b> .38	08.1		34	•
\$ \$	1409509	AAB68929	28.2	0.75	3.78	0.36	5.86	11.75	80.0		
; ;	365707	AA025434	3	0.53	3.68	0.58	3.38	10.11	1.36	9 6	٦ ،
3 8	750057	AA424920	£7 £	9	3.66	0.95	5.41	10.03	0.93	78.0	4 (
3 \$	16763B	W94121	80	0.49	3,65	5.62	0.55	0.68	2.00	4.9	<b>v</b>
5	30.00		}	:	† !	-					

Table 2-7; Good vs. Poor Clinical Outcome. Genes upregulated 3 fold or more in at least 1 tumor sample.

	7	7	7	•		4 (	7 (	2	N	7	8	,	4 (	4 (	.4	7	8	8	64	7	7	~	, ,	4 (	7	7	7	7	7	7	7	7	2	?	7	7	7	2	7	~	~	7	7	7	7	~	7	~		
	3.86	1.24	433	90	20.	4	3,45	0.90	1.8.	4.40	90		1.32	5.80	4.05	4.08	4.96	3.89	1.53	3.95	0.97	. C. E.	20.0	4.5	4.38	3.80	1.03	3.61	33.34	3.05	4.26	4.08	3.30	3.07	3.11	3.16	3.85	4,12	4.24	3.47	3.05	0.61	3.76	3.15	1.1	0.61	0.49	0.48	•	
	2.42	344	900	9 9	3.8C	0.18	4. 0	0.74	6,63	0.24		5 6	3.83	0.51	5.78	5.39	0.23	9	7.	. p	9 9	2 5	1.6/	2.03	3.46	1.07	1.06	0.44	28.	2.36	0.40	4.51	9 4	50 6	0.93	3.41	3.03	0.54	0.58	3.49	0,41	0.75	3.65	3.37	3.19	0.56	0.63	7	•	
	97.0	· •	34.	3	<u>.</u>	90.0	0.30	3.34		7.0	C. 7	0.91	2.15	0.45	0.43	0.69	0.25		- 6	S (5	9 ·	3.40	3.00	0.93	1.22	0.64	3.02	1 80	0.49	27.0	; <b>;</b>	5 5	7.00		65.0	8 6	2,00	. 750	80.0	0.85	1.87	4 05	5.50		3.72	27.5			0.4.0	
	Ş	76.	Ç.	4.0	1.55	2,10	0.65	200	. c	0.83	5.59	1.01	0.82	0.78	5.5	6.00	7 7 7	÷ 6	) (S	9. c	0.49	3.05	5.1	050	290	5	6	76.1 C3.C	3 4	8 6	<b>S S</b>	2.5	S (	60.	5. 5 6. 6		0.83	2.0	, c.	20.0	8.0 E		25.50	9.5	24.0			o. Jo	0.93	
	,	3.36	3.94	3.32	3.12	4.65	284	7.0.7	رار ر د د د	3.34	0.76	0.58	3.06	9 6	2 6	- C	7. 7.	1.03	3.28	4.01	9,34	1.59	0.89	3.85	0.75	2 6	8 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6 6	20.0	48.0	3.08	1.09	0.56	0.27	0.76	1.43	3.13	1.39	1.40	0.92	62.5	 4	16.0	0.55 6.6	0.42	0.69	0.26	0.57	0.36	9.0	က
		2.36	2.32	2.31	230	2.5	77.7	2.20	2.23	2.23	2.22	222	;	0.40	81.7 61.0	2.19	2.18	217	2.17	2.14	2.13	2.11	2.4	- <del>-</del> -		2.10	5.09	2.07	2.06	5.06	2.04	2.01	1.99	1.99	1.98	1.97	1.96	1.96	1.56	4.89	4.89	- 89	1.86	1.80	1.7	1:1	1.76	1.69	1.66	
		1.33	0.33	2	75.0	3 5	0.63	1.26	חיק.	2.14	12.67	2	ţ, Ş	1.40	17.15	15.57	13.30	2.98	0.24	1.57	10.50	17.		S :	16.27	0.86	0.30	0.83	10.94	9.27	3.81	0.01	25.44	2.07	1.1	1.50	0.85	1.51	2.19	8.31	0.76	15.04	13.84 48.64	3.24	21.21	27.34	3.24	9.0	0.02	
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2.43		70.7	0.81	66.0	5.19	<u>.</u>	3.60	1.47	2.04	2.98	1.75	1.38	1.53	3.29	3.41	1.13	6.42	5.23	1.57	.03 50.1	4.	3.56	1.81	1.26	0.90	1.7	1.80	0.98	3.17	3.19	2.17	0.85	7. S		17.7	 	50.5 70.4	85.0	3.44	; ;	ò	5 6	9.7	2.42	6.84	œ
204	9 6	2.01	2.01	2.01	2.00	2.00	2.00	2.00	1.99	1.99	1,99	1,99	1.98	1.98	1.98	1.98	1.97	1.97	1.97	1.97	1.96	1.96	1.96	1,96	1,96	1.95	1.85	1.95	1.95	1.94	<b>2</b> .	1.94	<del>.</del> .	3. 7	<b>3</b> . 3	<b>.</b> .	Se. →	1.93	8: +	8.	SB. 1	.93	.93	1.93	1.93	
,	5.6	2.33	0.92	0.62	0.77	0.48	11.14	3.88	9	10.17	0.35	0.36	1.22	3.01	2.09	0.48	3.29	1.52	2.32	3.21	0.28	6.16	2.48	1.06	1.24	0.01	1.19	2.47	7.09	0.43	277	<u>5</u> .	5.12	0.33	1,13		7.7	9.5 5.5	7.0	5.08 2.08	Z	9.08	4.49	20.	1.58	
3	8.52	4.69	1.85	1.24	 2	0.92	22.32	7.73	0.09	20.24	0.70	0.72	2.42	5.97	4.15	0.95	6.49	2.99	4.57	6.31	5.5	12.08	4.83	2.08	2.42	0.02	231	4.82	13.82	0.85	4.32	2.02	9. 2.	<b>2</b>	2.19	0.22	 	<u>.</u> .	0.72	5.76 5.36	4	0.12	8.66 8.66	1.24	3.05	
A 4 700 4 40		H38650	W15263	H84369	N38860	AA633866	AA630771	H84871	H15417	H77297	AA404293	AA464935	AA017133	AA421270	770892	W86987	AA453015	H14057	N89676	AA486435	R42520	H42679	R96827	AA427740	AA018460	AA443649	R34331	AA453997	AA962236	H23137	AA009609	AA128407	AA055835	AA707872	AA004835	104216	VV80055	A A 702207	100000 A	A4125825	H18423	AA777700	AA121836	R08266	N91914	
0,000	910000	190732	322537	249526	279905	858188	856420	249603	49333	233464	758366	838899	362628	731031	83999	416556	788334	163841	307053	842918	29954	183337	200741	770837	362424	771258	136474	785262	1802120	52303	365515	626587	377461	413150	429148	80226	416833	132954	450050	548693	50877	449257	566106	127193	306829	4
ě	Š	8	330	3	312	313	314	315	316	347	348	319	33	3 5	32	S	3	325	328	3.	£ 25	2 2		Ę	3 2	S	2	25	8	33,	88	88	Z	ੜ	8	<b>X</b>	<b>X</b>	£ :	<b>3</b>	*	<b>8</b>	ž	38	351	352	

Table 2-7: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

1572599   147449   245   1459   1459   159   150   1		900									•
AACTSTATE 1.55 0.187 1.55 0.70 0.82 0.58 1.25 0.58 0.58 0.58 0.58 0.58 0.58 0.58 0.5	,	60.	6	8	183	0.94	0.71	5.07	1.97	4 80 E	
AAGSTASS 1.57 1.47 1.52 5.76 0.82 0.558 1.42  AAGSTASS 7.48 0.89 1.22 3.57 1.62 0.59 1.44  AAGSTASS 7.48 0.89 1.82 3.57 0.94 0.94 1.24  AAGSTASS 7.48 0.89 1.82 3.57 0.94 0.94 1.24  AAGSTASS 7.48 0.89 1.82 3.55 1.77 1.62 1.58  AAGSTASS 7.48 0.89 1.82 2.53 3.56 1.77 1.62 1.58  AAGSTASS 7.48 0.89 1.82 2.53 3.56 1.77 1.62 1.58  AAGSTASS 7.48 0.89 1.82 0.85 1.92 0.87 1.77 1.62  AAGSTASS 7.48 0.89 1.82 0.85 1.92 0.87 1.77 1.62  AAGSTASS 7.48 0.89 1.82 0.85 1.95 1.77 1.62  AAGSTASS 7.49 1.45 1.89 1.89 0.87 1.77 2.64 1.97 0.88  AAGSTASS 7.41 0.80 1.89 1.80 0.87 1.77 1.46  AATTACL 1.50 1.89 1.89 1.80 0.87 1.77 1.46  AATTACL 1.50 1.89 1.89 1.89 1.89 1.89 1.89 1.89 1.89	<b>2</b> 2 (2	141408	200	 	193	1.90	1.30	1.58	92.7	2.5	-
AAASSSATI         1.10         1.45         1.52         3.51         1.16         2.69         1.40           AAAZSSATI         1.53         1.45         1.52         3.57         1.16         2.69         2.20           AAAZSSATI         1.53         0.80         1.52         2.37         0.94         0.94         2.41           AAASSSATI         1.53         0.86         1.92         2.35         0.77         1.91         2.41           AAASSSATI         0.16         1.92         2.35         0.77         1.91         1.37           AAASSSATI         0.16         0.15         1.92         0.84         1.24         1.38           AAASSATAS         0.15         0.24         1.92         0.84         1.27         1.97         1.97           AAASSAGAS         1.16         0.16         0.17         1.17         1.66         1.17         1.18         1.11         1.13         1.11         1.13         1.11         1.11         1.13         1.11         1.13         1.11         1.13         1.11         1.11         1.11         1.12         1.11         1.11         1.11         1.11         1.11         1.11         1.11         1.	88	AMB63/33	1.51	5 ,	2	5.76	0.82	0.58	C7'L	1 0	-
AAA28117 1,80 0.94 132 3.7 0.94 0.99 0.20  AAA21728 7.4 3.89 132 3.5 0.94 0.44 2.41  AA686418 0.99 0.05 1122 0.35 0.77 162  AA46087 0.03 0.15 1122 0.35 0.97 0.94 0.41  AA46087 0.03 0.15 1122 0.35 0.97 1.05  AA46087 0.03 0.15 1122 0.35 0.97 1.05  AA46087 0.03 0.15 1122 0.35 0.97 1.05  AA46087 0.03 0.03 1.02 0.03 0.04 0.07  AA46087 0.03 0.03 1.02 0.03 0.07  AA46087 0.03 0.03 0.03 0.03 0.03  AA6657 0.03  AA6657 0.03 0.03  AA6657 0.03  A	છ	AA953973	279		76.1	2 54	1 16	2.69	1.40	8	
AASTSSTO         15.3         0.80         1.92         2.57         0.78         0.41         2.44           AATTOSOT         0.05         0.05         1.92         2.53         3.56         1.34         1.38           AATSOT         0.05         0.05         1.92         2.53         3.56         1.34         1.38           AATSOT         0.05         0.05         1.92         2.53         1.74         1.01         0.75           AAATSATAS         0.05         0.05         1.92         2.23         3.05         0.07         1.37           AAATSATAS         0.05         0.05         1.92         2.23         3.05         0.07         1.37           AAATSATAS         0.05         0.05         1.24         1.24         1.01         0.75         1.07         1.05         0.07         1.07         0.07         1.07         0.07         1.05         1.15	10	AA426113	1.80	<b>9</b> .0	1.92	9.0	70	0.99	2.30	2.04	
AMSTORE         7.48         3.89         1.92         2.35         3.69         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.39         2.35         1.77         1.62         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.38         1.39         1.38         1.39         1.38         1.39         1.38         1.39         1.38         1.39         1.38         1.39	~	AA235370	1.53	0.80	1.92	, c, c	2 C	0.41	2.41	2.47	<b>-</b> - ·
AA419077         0.09         0.05         1.92         2.53         1.77         1.59         1.55         1.75         1.55         1.15         1.55         1.15         1.55         1.15         1.55         1.15	_	AA670286	7.48	3.89	1.92	S. 30	) c	7	1.38	1.02	_
AA60827         0.35         0.16         1.92         3.35         1.77         1.99         3.12         1.97         3.12         1.97         3.12         1.97         3.12         1.97         3.12         1.97         3.12         1.97         3.12         4.00		AA130017	0.09	0.05	1.92	3.7	3 5	167	1,36	1.51	-
AGESTALE         0.86         0.51         1.92         0.84         0.93         1.92         0.84         0.93         1.92         0.94         0.93         1.92         0.94         0.93         1.92         0.94         0.93         1.92         0.84         0.93         1.15         1.11         1.14         1.15         1.11         1.15		AA460827	0.35	0.18	1.92	3.35		3 52	1.97	2.63	-
A0085748         11.69         6.09         1.92         0.85         USF         <	, ~	AA666418	0.98	0.51	1.92	9. 26.	<b>5</b> 6		3.54	2.37	-
Mail		AA085748	62	609	1.92	0.85	76.0	) (1)	. c	1.12	-
AAA78456         16.65         8.71         1.92         2.29         3.06         0.06/1         1.10           AAA8842         1.65         8.71         1.91         3.07         1.56         1.41         1.15		TA0005	28.0	0.43	1.92	5.44	1.24	1.0	, ,	187	-
AAA97265         1.51         1.59         1.31         0.42         2.10           AAA97265         1.56         1.51         1.59         1.31         0.42         2.11           AAA9362         1.23         6.47         1.91         3.62         1.55         1.41         1.15           AAA9362         1.26         1.91         3.70         2.54         1.97         0.68           AAA93627         2.20.4         11.54         1.91         3.70         2.54         1.97         0.68           AAA93637         1.22         0.64         1.90         2.47         0.68         0.57         0.67         0.69           AA37649         1.22         0.64         1.90         0.77         0.89         0.67         0.89           AA37649         0.50         0.27         0.67         0.61         1.77         1.48         1.17         1.48         1.17         1.48         1.17         1.48         1.13         0.80         0.87         1.13         0.89         0.87         1.14         1.15         0.89         0.87         1.14         1.14         1.15         0.89         0.87         1.14         1.15         0.89         0.87		00000	7 7	9 6	1 92	2.29	3.06	. 79.0	2 9		-
AAATSAS         18 bbs         6.77         1.91         3.62         1.55         1.94         1.13           N91003         5.79         3.02         1.91         3.62         1.55         1.41         1.15           AABSTS 7         2.70         3.02         1.91         3.01         2.54         1.97         0.68           AABSTS 7         2.70         1.15-4         1.91         3.70         2.54         1.97         0.69           NAF3T 1-14         0.75         1.15         1.17         2.64         1.07         0.67         0.69           AAATSASS 2.22         1.15         1.19         3.77         2.64         1.07         0.67         0.68           AAATSASS 2.22         1.37         1.90         0.77         0.61         1.37         1.48         0.67         0.67         0.68         0.67         0.68         0.67         0.68         0.67         0.68         0.67         0.67         0.68         0.67         1.44         1.15         0.68         0.67         0.68         0.67         0.68         0.67         0.67         0.68         0.67         0.68         0.67         0.68         0.68         0.77         0.74 <t< td=""><td></td><td>790901</td><td>5.5</td><td>9 6</td><td>10.</td><td>190</td><td>1.31</td><td>0.42</td><td>2.10 2.</td><td>9 6</td><td>-</td></t<>		790901	5.5	9 6	10.	190	1.31	0.42	2.10 2.	9 6	-
NASO12   12.30   6.47   1.91   3.01   1.15	ເກ	AA487465	16.66	Z !		3.63	1.55	<b>1</b> .94	1.15		- 🕶
AdoS5768         176.33         3.02         181         3.01         2.54         1.97         0.68           AAOS5768         176.33         1.24         1.91         3.01         2.54         1.97         0.68           AAS568         1.22         0.64         1.90         0.74         3.82         1.07         0.57           H7406         1.22         0.64         1.90         0.74         3.82         1.07         0.57           AA35695         4.11         2.17         1.90         0.74         3.82         1.07         0.57           AA35695         4.11         2.17         1.90         0.71         0.89         1.07         1.48           AA5004         2.52         1.33         1.90         0.71         0.89         1.17         1.48         1.30         0.87         1.17         1.48         1.17         1.48         1.17         1.48         1.17         1.48         1.17         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14         1.14	~	N50912	12.38	6.47	1.91	2,0%		1.41	1.15	2.43	
AAOSSYGE         176.33         92.35         1.91         3.70         2.54         0.75         0.93           H74086         1.20         1.54         1.91         1.77         2.64         0.75         0.81         0.77         0.81         0.75         0.89           NZ8771         1.42         0.76         1.90         0.77         0.81         0.77         0.80         0.77         0.80         0.77         0.80         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.77         0.89         0.87         0.89         0.87         0.89         0.87         0.89         0.87         0.89         0.87         0.89         0.87         0.89         0.87         0.89         0.87         0.89         0.89         0.87         0.89         0.87         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89         0.89	80	N91003	5.78	3.02	1.91	3.0.	3.5	197	0.68	0.65	-
AA68267         22.04         11.54         1.91         1.77         2.54         1.07         0.57           H74086         1.22         0.64         1.90         2.47         3.82         1.07         0.57           N35175         1.45         0.76         1.90         2.47         3.82         1.07         0.57           AA158695         4.11         2.17         1.90         0.71         0.61         6.72         0.89           AA55869         4.11         2.17         1.90         0.71         0.61         1.17         1.46           AA560492         2.22         1.30         1.90         1.16         1.17         1.46           AA504492         2.77         1.69         2.47         3.61         1.31         1.44         1.17         1.46           AA504492         2.77         1.69         2.72         1.49         1.69         3.16         1.31         1.44         1.17         1.46         1.17         1.46         1.17         1.46         1.17         1.46         1.17         1.46         1.17         1.14         1.17         1.14         1.17         1.14         1.17         1.17         1.17         1.14		AA055768	176.33	92.35	1.91	3.70	<b>7</b> .7	42.0	0.93	3.44	-
HY4086 1.22 0.64 1.90 2.47 3.82 1.00 0.80  ANSESTS 4.11 2.17 1.90 0.71 0.81 6.72 0.80  ANSTAGO 2.24 1.90 0.71 0.81 6.72 0.80  ANSTAGO 2.24 1.90 0.71 0.81 1.16 1.31  N7301 2.84 1.50 1.89 1.00 1.16 1.31  AAGOMAG 2.77 1.46 1.89 1.00 1.80 1.80 0.85  N88576 0.50 0.77 1.89 1.60 1.80 0.85  AASSAGO 3.08 1.63 1.89 1.89 0.89 1.80 0.86  AASSAGO 3.08 0.70 1.88 0.70 1.48 1.89 1.42 1.88 1.44  AASSAGO 3.28 1.71 1.88 0.72 1.89 1.89 0.80  AASSAGO 3.28 1.72 1.72 1.89 1.89 0.80  AASSAGO 3.28 1.73 1.87 1.72 1.89 0.80  AASSAGO 3.28 1.75 1.87 1.72 1.89 0.80  AASSAGO 3.38 1.81 1.81 1.80 0.71 1.70  AASSAGO 3.38 1.81 1.81 1.80 0.71 1.70  AASSAGO 3.38 1.81 1.81 1.80 0.71 1.70  AASSAGO 3.38 1.81 1.81 1.80 1.71 1.72 1.89 0.89  AASSAGO 3.38 1.81 1.81 1.80 1.71 1.72 1.89 0.89  AASSAGO 3.38 1.81 1.81 1.80 1.81 1.80 0.89  AASSAGO 3.38 1.81 1.88 2.83 0.39 1.87 1.72 1.89 0.89  AASSAGO 3.38 1.81 1.88 2.83 1.85 1.85 0.89  AASSAGO 3.38 1.81 1.88 2.83 1.83 1.83 0.89  AANTORAO 1.72 0.83 1.85 1.85 1.89 0.89  AANTORAO 1.72 0.83 1.85 1.85 1.70 0.89 1.85  AANTORAO 1.72 0.83 1.85 1.85 1.70 0.89 0.89  AANTORAO 1.72 0.89 1.85 1.85 1.70 0.89 0.89  AANTORAO 1.72 0.89 1.85 1.85 1.70 0.89 0.89  AANTORAO 1.72 0.89 0.89 0.80 0.80 0.80 0.80 0.80 0.80	<b>,</b> 4	AA687677	22.04	7	1.91	1.77	2.64	9 7	75.0	1.59	-
NATOR   1.22   1.50   0.77   1.60   0.74   0.81   0.75   1.60   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75   1.50   0.75		72708	7	0.54	1.90	2.47	3.82	7.07		0.46	-
AA55556         4.17         1.45         1.17         1.45           AA57556         2.52         1.37         1.90         3.33         1.64         1.17         1.40           AA57556         2.52         1.37         1.90         1.46         1.89         2.02         1.49         3.16         0.87           AA57556         2.52         1.37         1.90         1.76         1.46         3.16         0.87           AA50492         2.77         1.46         1.89         4.59         1.00         1.87         1.04           AA7643         0.50         0.27         1.89         1.69         3.23         1.87         1.05           N6876         0.12         0.07         1.89         4.90         1.80         0.87         1.89         1.89         1.89         1.89         0.85         0.87         1.89         0.85         1.18         0.85         0.87         1.89         0.85         1.18         0.85         0.85         0.85         0.87         1.89         0.85         1.18         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85         0.85 <td< td=""><td>20 1</td><td>2004</td><td>77.</td><td>9 6</td><td>1 90</td><td>0.71</td><td>0.81</td><td>6.72</td><td>8.9</td><td>4 B7</td><td>-</td></td<>	20 1	2004	77.	9 6	1 90	0.71	0.81	6.72	8.9	4 B7	-
AASTSSSS         4.11         2.17         1.50         1.15         3.16         1.31           AASTSSSS         2.52         1.53         1.69         1.69         1.69         3.16         1.31           AASTSSSS         2.52         1.50         1.89         2.02         1.49         3.16         1.31           AASTSSS         2.57         1.46         1.89         2.02         1.77         5.41         1.04           AASTSAGS         0.17         1.89         1.69         1.00         1.86         0.87           NGSTF         0.17         1.89         1.69         1.80         0.87         1.89         1.69         0.87           AASSBS         1.51         0.80         1.89         4.90         1.80         0.87         1.89         1.80         0.87         1.89         1.84         1.89         1.89         1.84         1.84         1.89         1.84         1.89         1.89         1.89         1.84         1.89         1.84         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89         1.89	S.	1/197N	45	9 1	200	333	<u>-</u> . 2	1.17	<del>2</del>	0.	•
AA575556         2.52         1.33         1.31           AA575576         2.52         1.49         3.16         1.31           AA50442         2.77         1.46         1.89         4.59         1.00         1.66         0.87           AA776434         2.77         1.46         1.89         4.59         1.00         1.66         0.87           AA776434         0.50         0.27         1.89         1.69         3.23         1.87         1.50           N68576         0.12         0.07         1.89         1.69         3.23         1.87         1.50           N68576         0.12         0.07         1.89         1.69         3.23         1.87         1.50           AA38924         1.51         0.80         1.89         5.55         1.18         0.80         0.87           AA38925         1.51         0.80         1.88         0.93         1.18         0.41         0.41           AA38926         1.51         0.80         1.88         0.93         1.18         0.41         0.41         0.41         0.41         0.41         0.41         0.41         0.41         0.41         0.41         0.41         0.41	<b>⊡</b>	AA155695	4.11	2.1/	) (c	1 16	1,15	3.09	2.42	66.	
AA73301         2.84         1,50         1,89         4,50         1,00         1,86         0.87           AA776434         0.50         0.27         1,89         1,02         1,17         5,41         1,04           AA776434         0.50         0.27         1,89         1,69         1,17         5,41         1,04           N68576         0.12         0.07         1,89         1,69         1,18         0.87         1,89         1,69         1,18         0.75           AA393408         1,51         0.80         1,89         5,55         1,18         0.66         0.73           AA393408         1,51         0.80         1,89         0.55         1,18         0.66         0.73           AA393408         1,51         0.80         1,88         0.93         1,18         0.66         0.73           AA393408         1,51         0.80         1,88         0.93         1,18         0.66         0.73           R53967         0.76         1,88         0.06         1,44         0.44         0.44         0.44           AA39408         1,51         0.80         1,88         0.05         1,16         1,44         0.44	51	AA975556	2.52	1.33	06.1	50.0	1.49	3.16	1,31	, t	- •
AA50442         2.77         1,46         1,89         1,72         1,17         5,41         1,04           AA776434         0.50         0.27         1,89         1,69         1,89         1,69         1,17         5,41         1,04           No8376         0.12         0.07         1,89         1,69         1,18         0,66         0,73           H09936         3.08         1,83         4,90         1,80         0,85         0,87         0,87         0,87           AA393408         1,51         0.80         1,88         0,93         1,18         0,47         1,54	<b>8</b> 0	N73301	2. <b>8</b> 4	1.50	20°C	<b>2</b> .02	60	1.86	0.87	1.14	- •
AA776434         0.50         0.27         1.89         1.02         3.23         1.87         1.50           N68576         0.12         0.07         1.89         1.69         3.23         1.87         1.50           H09356         0.12         0.07         1.89         4.90         1.80         0.86         0.07           AA389408         1.51         0.80         1.89         4.90         1.80         0.73           AA389408         1.51         0.80         1.89         0.93         1.18         0.66         0.77           R53963         0.76         0.40         1.88         0.93         1.18         0.41         0.47           R53963         0.76         0.40         1.88         0.95         1.18         0.47         1.44           R53963         0.76         0.04         1.88         0.06         0.71         0.47         1.44         0.47           R63964         0.51         1.88         0.02         1.88         0.02         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05 <td< td=""><td>φ</td><td>AA504492</td><td>2.77</td><td>1.46</td><td>1.89</td><td>8.C.</td><td>1 17</td><td>5.41</td><td><b>2</b>.</td><td>0.81</td><td>- ,</td></td<>	φ	AA504492	2.77	1.46	1.89	8.C.	1 17	5.41	<b>2</b> .	0.81	- ,
N68576         0.12         0.07         1.89         1.69         0.20         0.86         0.07           AA383408         1.51         0.80         1.63         4.90         1.89         0.66         0.73           AA383408         1.51         0.80         1.63         1.89         4.90         0.66         0.73           AA383408         1.51         0.80         1.88         0.93         1.83         3.74         1.54           AA38922         11.03         5.86         1.88         0.93         1.83         3.74         1.54           AA38922         11.03         0.86         1.69         1.69         0.66         0.47         0.47           R53963         0.76         0.92         1.88         3.06         1.13         1.64         1.44           R05416         0.04         0.02         1.88         3.06         1.72         1.28         0.80         1.05           H1176         0.08         0.52         1.88         0.72         1.23         6.60         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05         1.05	φ	AA776434	0.50	0.27	1.89	7.02		1.87	5.50	1.16	_
H09936         3.08         1.63         1.89         4.90         1.80         0.03         0.73           AA393408         1.51         0.80         1.89         4.90         1.80         0.05         0.73           AA39802         1.51         0.80         1.89         4.90         1.69         1.18         0.04         1.54	9	N68576	0.12	0.07	1.89	1.69	2.5	. c	0.87	1.00	-
AA398408         1.51         0.80         1.89         5.55         1.18         0.00         0.00           AA398922         11.03         5.86         1.88         0.93         1.18         0.00         0.47         1.54         0.47         1.54         0.47         1.54         0.47         1.54         0.47         1.54         0.47         0.47         1.54         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44         0.47         0.44	? a	H09936	3.08	1.83	1.89	4.90	08.1	9 6	27.0	1.30	-
AA398922         1.51         0.93         1.83         3.74         1.57           R53983         1.03         5.86         1.88         0.93         1.83         3.74         1.57           R53983         0.76         0.40         1.88         3.06         1.42         1.28         0.80           R53963         0.76         0.80         1.88         3.06         1.42         1.28         0.80           R05416         0.04         0.02         1.88         0.72         1.23         6.60         1.05           H11376         0.98         0.52         1.88         0.72         1.23         6.60         1.05           AA088517         2.26         1.21         1.87         1.72         1.69         3.24         1.05           AA77659         3.28         1.75         1.87         1.77         1.09         1.77         1.09           AA77649         0.56         0.49         1.87         1.87         1.27         3.59         1.07         1.07           AA77647         2.26         1.87         1.87         1.27         1.09         1.17         1.07           AA927125         3.38         1.81 <t< td=""><td>o \$</td><td>A A 202408</td><td></td><td>O BC</td><td>1.89</td><td>5.55</td><td>1.18</td><td>0.00</td><td>2 7</td><td>1.37</td><td>-</td></t<>	o \$	A A 202408		O BC	1.89	5.55	1.18	0.00	2 7	1.37	-
AA39632         11.03         1.69         1.16         1.16         0.41         0.41           N63363         0.76         0.30         1.86         3.06         1.13         1.86         0.40           N63361         1.51         0.80         1.88         3.06         1.13         1.26         0.80           R05416         0.04         0.02         1.88         3.00         1.42         1.28         0.80           H11376         0.98         0.52         1.88         0.72         1.23         6.60         1.05           AA757659         3.28         1.75         1.87         4.39         2.04         0.91         0.75           AA757659         3.28         1.75         1.87         1.77         1.02         1.07           AA757659         3.28         1.75         1.87         3.46         1.69         1.77         1.09           AA757659         3.28         1.87         3.46         1.69         1.77         1.07         1.07           AA757659         3.28         1.87         5.34         0.66         0.70         1.07           AA75747         22.84         1.2.15         1.86         2.37	2	000000	<u> </u>	9	1 88	0.93	.8. 8.	3,74	<u>,                                    </u>	. H	-
N64395         1.51         0.80         1.88         3.06         1.13         1.86         1.44           N64391         1.51         0.80         1.88         3.06         1.42         1.28         0.80           R05416         0.04         0.02         1.88         3.00         1.42         1.28         0.80           H1376         0.96         0.02         1.87         4.39         2.04         0.91         0.75           AA757659         3.28         1.75         1.87         4.39         2.04         0.91         0.75           AA757659         3.28         1.75         1.87         1.77         1.02         0.75         1.02           AA757659         3.28         1.75         1.87         3.46         1.60         1.77         1.02           AA286814         0.58         0.30         1.87         3.46         1.60         1.77         1.02           AA286814         0.58         0.09         1.87         3.46         1.60         1.77         1.07           AA286814         0.58         0.09         1.87         0.49         1.87         0.31         0.32           AA77547         22.84	<b>X</b> :	AA398922	3.5	9 6	88 -	1.69	1.18	0.41	, C. 4.	5 -	_
NOSASSI         1,51         0.00         1,88         3.00         1,42         1,28         0.80           H1376         0.04         0.02         1,88         3.00         1,42         1,28         0.80           H1376         0.98         0.52         1,88         0.72         1,23         6.60         1.05           AA75769         3.26         1,21         1,87         1,77         1,69         3.24         1,02           AA75769         3.28         1,75         1,87         1,77         1,69         3.24         1,02           AA75769         3.28         1,87         3.46         1,60         1,77         1,02           AA75769         3.28         0.30         1,87         3.46         0.66         0.70         1,07           AA77547         22.84         1.87         5.34         0.66         0.70         1,17         1,09           AA77547         22.84         1.2.25         1.86         2.37         0.83         0.31         3.50           AA77547         22.84         1.2.1         1.86         5.12         1.28         0.88         1.12           AA37755         3.38         1.81	<b>n</b>	200207	0.5	9 6	1 88	3.06	1,13	1.85	<b>\$</b> . \$	 	-
R03410         0.04         0.05         1.88         0.72         1.23         6.60         1.05           AA086517         2.26         1.21         1.87         4.39         2.04         0.91         0.75           AA757659         3.28         1.75         1.87         1.72         1.69         3.24         1.02           AA286814         0.56         0.30         1.87         3.46         0.66         0.70         1.07           AA757659         3.28         1.87         3.46         1.60         1.77         1.02           AA286814         0.56         0.30         1.87         3.46         0.66         0.70         1.07           R91220         0.92         0.49         1.87         1.90         1.27         5.59         0.92           AA77547         22.84         12.25         1.86         2.37         0.63         0.51         1.12           AA77547         22.84         1.25         1.86         5.12         1.28         0.58           AA77547         22.84         1.25         1.86         5.17         0.59         1.52           AA351725         3.38         1.86         1.16         1.16	25	07702		9 6	1 88	3.00	1,42	1.28	G.80	, c	-
AAR7355 61.64 33.03 1.87 4.39 2.04 0.91 0.75 AAR75759 3.28 1.27 1.87 1.72 1.69 3.24 1.02 AAR75759 3.28 1.37 1.87 1.72 1.69 3.24 1.02 AAR7355 61.64 33.03 1.87 5.34 0.66 0.70 1.07 AAR7355 61.64 33.03 1.87 5.34 0.66 0.70 1.07 AAR7355 61.64 33.03 1.87 1.00 1.27 5.59 0.82 AA77547 22.84 12.25 1.86 5.12 1.28 0.89 1.12 AA77547 22.84 12.25 1.86 5.12 1.28 0.89 1.52 AA87355 0.02 0.01 1.86 1.18 0.08 4.59 1.52 AA873515 4.89 2.63 1.81 1.86 1.16 3.75 0.99 2.02 AA87515 0.02 0.01 1.85 1.16 0.65 0.65 AA777840 1.72 0.93 1.85 1.70 1.31 3.01 0.97 AA707840 1.72 0.93 1.85 1.36 1.36 1.36 1.58 AA707840 1.72 0.93 1.85 1.36 1.36 1.36 1.36	¥	703410	\$ 5 5 6	4 6	88	0.72	<u>5</u>	9.60	D		-
AA088317         2.26         1.21         1.69         3.24         1.02           AA757659         3.28         1.75         1.87         1.77         1.09           AA28814         0.58         0.30         1.87         3.46         0.66         0.70         1.07           AA37355         61.64         33.03         1.87         1.00         1.27         5.59         0.92           AA77547         22.84         12.25         1.86         2.37         0.83         1.12         0.88         1.12           AA77547         22.84         12.25         1.86         5.12         0.83         0.31         1.12           AA77547         22.84         12.25         1.86         5.12         0.83         1.52         0.88         1.12           AA7547         22.84         1.26         0.83         1.86         1.18         0.89         4.59         1.52           AA931725         3.38         1.81         1.86         1.16         3.75         0.89         1.53           AA52361         0.02         0.01         1.86         1.16         3.75         0.89         1.54         3.32           AA707840         1.72 </td <td>9</td> <td>H113/6</td> <td>0.80</td> <td>70.0</td> <td>187</td> <td>4.39</td> <td>2.04</td> <td>0.91</td> <td>0.73</td> <td>07.</td> <td>-</td>	9	H113/6	0.80	70.0	187	4.39	2.04	0.91	0.73	07.	-
AA757659         3.28         1.79         1.09           AA28684         0.58         0.30         1.87         3.46         1.60         1.77         1.09           AA28684         0.58         0.30         1.87         3.46         0.66         0.70         1.07           AA37355         61.64         33.03         1.87         1.00         1.27         5.59         0.92           AA77547         22.84         12.25         1.86         2.37         0.63         0.31         3.50           AA77547         22.84         1.25         1.86         5.12         1.28         0.89         1.52           AA7547         22.84         1.25         1.86         5.12         0.89         4.59         1.52           AA3535         0.32         1.86         1.18         0.89         4.59         1.53           AA707840         1.72         0.93         1.85         2.40         0.65         1.54         3.32           AA707840         1.72         0.93         1.85         1.70         0.57         1.54         3.32           AA707840         1.72         0.93         1.85         4.35         1.36         1.58	33	AA088517	2.26	17.1	10.4	1.72	1.69	3.24	1.02	9.	٠ -
AA286814         0.58         0.30         1.07         1.07           R91220         0.92         0.49         1.87         5.34         0.66         0.70         1.07           AA873355         61.64         33.03         1.87         1.00         1.27         5.59         0.62           AA873355         61.64         33.03         1.87         1.00         1.27         5.59         0.62           AA93175         2.84         1.21         1.86         5.12         1.28         0.89         1.12           AA93175         3.38         1.81         1.86         5.12         0.89         4.59         1.52           AA30175         3.38         1.81         1.86         1.16         3.75         0.99         2.02           AA526315         4.89         2.63         1.86         1.16         3.75         0.99         2.02           H52361         0.02         0.01         1.86         1.76         0.65         1.54         3.32           AA707840         1.72         0.93         1.85         1.70         1.31         3.01         0.97           R26859         1.06         0.57         1.85         4.35	45	AA757659	3.28	1.73		3.45	1.60	1.1	1.09	\$ 5	- •
R912ZO         0.92         1.07         1.27         5.59         0.92           AA873355         61.64         33.03         1.87         1.00         1.27         5.59         0.92           AA77547         22.84         12.25         1.86         2.37         0.83         1.12           AA931725         3.38         1.81         1.86         5.12         1.28         0.89         1.52           AA626315         4.89         2.83         1.86         3.17         1.07         0.72         1.83           H52351         0.02         0.01         1.86         1.16         3.75         0.99         2.02           AA707840         1.72         0.93         1.85         2.40         0.65         1.54         3.32           R26859         1.06         0.57         1.85         4.35         1.56         0.80	2	AA286814	0.58	0.30	.0.	2	0.66	0.70	1.07	 60.	- •
AA873355         61.64         33.03         1.01         2.37         0.63         0.31         3.50           AA77547         22.84         12.25         1.86         2.37         0.63         0.31         3.50           N92012         3.38         1.01         1.86         5.12         1.28         0.89         1.52           AA93125         3.38         1.81         1.86         1.18         0.69         2.02         1.83           AA626315         4.89         2.63         1.86         1.16         3.75         0.99         2.02           H52361         0.02         0.01         1.86         1.16         3.75         0.99         2.02           AA707840         1.72         0.93         1.85         1.70         1.31         3.01         0.97           R26859         1.06         0.57         1.85         4.35         1.36         1.58         0.80	21	R91220	0.92	0.49	1.0	5 5	1.27	5.59	0.92	2. 2. (	- •
AA775447 22.84 12.25 1.09 2.21 1.28 0.89 1.12 N92801 1.88 1.01 1.86 5.12 1.28 0.89 1.52 AA931725 3.38 1.81 1.86 1.18 0.89 4.59 1.52 AA626315 4.89 2.63 1.86 1.16 3.75 0.99 2.02 H52351 0.02 0.01 1.85 2.40 0.65 1.54 3.32 AA707840 1.72 0.93 1.85 1.70 1.31 3.01 0.97 R26859 1.06 0.57 1.85 4.35 1.36 1.58 0.80	ž	AA873355	61.64	33.03	20.7	75.6	0.83	0.31	3.50	2.52	- •
N92801         1.88         1.01         1.89         3.14         4.59         1.52           AA931725         3.38         1.81         1.86         1.18         0.69         4.59         1.52           AA26315         4.89         2.63         1.86         3.17         1.07         0.72         1.93           H52361         0.02         0.01         1.86         1.16         3.75         0.99         2.02           R43535         0.73         0.39         1.85         2.40         0.65         1.54         3.32           AA707840         1.72         0.93         1.85         1.70         1.31         3.01         0.97           R26859         1.06         0.57         1.85         4.35         1.36         1.58         0.80	82	AA775447	8	12.25	08.7	, , , , , , , , , , , , , , , , , , ,	1 28	0.88	1.12	0.93	
AA931725     3.38     1.81     1.85     1.19     0.72     1.93       AA626315     4.89     2.63     1.86     3.17     1.07     0.72     1.93       H52361     0.02     0.01     1.86     1.16     3.75     0.99     2.02       R43535     0.73     0.39     1.85     2.40     0.65     1.54     3.32       AA707840     1.72     0.93     1.85     1.70     1.31     3.01     0.97       R28659     1.06     0.57     1.85     4.35     1.36     1.58     0.80	9	N92801	1.88	1.01	99.	7 9	200	4,59	1.52	1.14	- •
AA626315         4.89         2.63         1.86         3.17         1.27         0.99         2.02           H52361         0.02         0.01         1.86         1.16         3.75         0.99         2.02           R43535         0.73         0.39         1.85         2.40         0.65         1.54         3.32           AA707840         1.72         0.93         1.85         1.70         1.31         3.01         0.97           R28859         1.06         0.57         1.85         4.35         1.36         1.58         0.80	88	AA931725	3.38	1.81	3.6	- C	\$ t	0.72	1.93	2.41	<b>-</b>
H52361 0.02 0.01 1.86 1.16 5.75 1.54 3.32 8.435 0.73 0.39 1.85 2.40 0.65 1.54 3.01 0.97 8.435 1.36 1.56 0.80 82859 1.06 0.57 1.85 4.35 1.36 1.56 0.80	2	AA626315	4.89	2.63	1.85		37.0	66.0	2.02	1,36	_
R43535 0,73 0.39 1.85 2.40 0.55 1.27 0.97  AA707840 1,72 0.93 1.85 1.70 1.31 3.01 0.97  R26859 1,08 0.57 1.85 4.35 1.36 1.58 0.80	<u>ر</u>	H52361	0.02	0.01	1.86	1.16	5.73	4.54	3.32	1.36	-
AA707840 1.72 0.93 1.85 1.70 1.31 5.01 0.57	ğ	R43535	0.73	0.39	1.85	2.40	0.65	<u> </u>	790	2.27	-
RZ6859 1.06 0.57 1.85 4.35 1.36 1.58 0.50	2 5	AA707840	2	0.93	1.85	1.70	1.31	3.U	6.0	1.18	-
	D 0	R76859	1 2	0.57	1.85	4.35	 8.	8c. L	0.0	•	
•	2		<u>:</u>			σ					

Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

1.25 3.19 0.97 0.95 0.86 1.48 1.17 1.17 0.63 0.88 1.44 1.44	1.38 1.37 1.37 1.37 1.37 1.20 1.20 1.39 1.09 1.09 1.03 1.03 1.03 1.03 1.03 1.03	
1.16 1.45 1.45 1.16 1.16 1.20 0.89 0.89 0.89 1.72 1.72 1.72 1.72 1.98 3.16	1,02 1,08 1,71 1,71 1,08 1,21 1,23 1,23 1,23 1,23 1,18 1,18 1,18 1,106 1,06 1,06 1,105 1,1	
5.04 4.17 7.14 4.60 9.99 9.16 9.18 1.89 1.89 1.89	1.07 4.94 4.13 3.09 4.14 0.65 3.26 4.97 4.97 1.11 1.11 1.37 1.37 1.37 1.37 1.37 1.3	
1.05 1.17 1.17 1.25 1.50 1.72 1.72 1.72 1.72 1.24 1.24 1.25 1.25 1.30	1.96 1.40 1.22 1.26 1.27 1.26 1.36 1.36 0.98 0.98 1.43 1.35 0.93 0.93 1.45 1.76 0.93 1.35 0.93 1.35 1.35 1.35 1.35 1.35 1.35 1.35 1.3	
0.78 1.440 1.440 1.02 1.02 1.03 1.03 1.88 1.88 1.88 1.86 1.86 1.86 1.12	3.69 1,10 1,10 1,03 1,04 1,104 1,105 1,106 1,106 1,106 1,106 1,106 1,107 1,106 1,107 1,106 1,107	<b>6</b>
1.85 1.85 1.85 1.84 1.84 1.84 1.83 1.83 1.83 1.83 1.83 1.83	1,82 1,82 1,82 1,82 1,82 1,82 1,81 1,80 1,80 1,80 1,80 1,80 1,80 1,80	
1.27 15.36 0.87 0.12 1.47 1.47 2.52 0.80 0.56 0.25 0.15 0.11 1.16 1.09	0.31 0.60 0.80 0.80 0.87 0.09 0.09 0.09 0.09 0.09 0.09 0.09 0.0	
2.36 28.41 1.51 0.21 2.35 1.19 2.71 4.64 1.03 0.21 1.30 1.30 2.00 3.46	0.66 0.06 0.06 0.00 1.45 4.49 0.17 1.58 1.58 1.72 1.72 1.72 1.08 0.70 0.80 0.80 0.80 0.80 0.80 0.80 0	
AA704587 AA886870 H22563 AA490987 H19825 AA493188 W32408 R80217 AA505045 H17696 AA411387 AA411387 AA411387 AA699582 AA099582	H10761 R48320 AA707935 AA098896 N47604 AA775863 AA872454 AA199075 AA458491 AA180204 AA775223 AA57223	
450598 1493252 51916 824573 172517 799878 321359 147050 825648 50043 754604 774078 1031278 51320 489706	47169 153541 392399 489553 277848 878544 1476019 645259 262060 812244 1555427 155862 200395 155817 461727 701112 1580342 46353 259884 1031598 271662 51773	
398 399 400 401 404 405 407 408 409 411 411 412 413 414 415 416	4	

. 543	757873	AA442853	7.87	4.41	1.78	1.42	1.72	1.06	4.23 0.94	0.50	
1 1	7,785	AA478479		5.50	1.78	1.44	4,45	0.80	1.41	0.80	
\$	245330	N54586	1.27	0.72	1.78	3.75	1.39	1.60	1.28	0.89 1	- •
4	731272	AA420993	7.07	3.98	1.78	3.50	0.87	1.30	1.65	1.5/ 28.	
844	1809036	A1000502	14.33	6.03	1.7	0.74	4.40	2 2	3.22	4.8	
449	451511	AA707336	0.68	0.38	1.77	1.20	4.74	1.29 22.6	80.0 •	7.0 97.6	- •-
450	39833	R53455	10.53	5.95	1.77	1.75	1.83	0.20	0 20 •	3.43	
451	21894	T66320	2.73	<u>.</u> %	1.77	1.17	1.91	0.50	. ·	24.8	
452	121948	T97762	9.91	5.60	1.72	0.47	1.82	1.12	70.	7 6	
453	343760	W69271	5.57	3.15	1.77	1.32	1.12	5.4.3		5. C	
454	51103	H19217	0.37	0.21	1.77	-30	0.85	4.47	<b>4</b> 8		
455	281936	N51069	1.68	0.95	1.71	3.4	20.	Ę.	8. 6	8 5	
<b>45</b>	562664	AA086204	1.47	0.83	1.76	1.18	1.98	3.74	E 6	5 5	•
157	248288	N78092	1.45	0.82	1.76	3.05	1.85	39.	2.5	2.5	
458	768130	AAA53310	118.87	67.49	1.76	2.31	2.20	3.20	0.55		. •
459	449033	AA777394	0.71	0.40	1.76	2.13	<u>2</u> .	0.83	<b>対</b> つ	07:1	- •
8	878231	AA775774	5,18	2.93	1.76	1.47	1.07	3.15	2.46	20.0	- •
461	138294	R56854	3.84	2.19	1.78	1.02	1.15	3.32	4.74	ָרָ מָלָ	- •
462	295843	N66957	4.79	2.73	1.76	0.55	0.61	0.35	1.52	07.0	- •
463	796287	AA460849	96.0	0.55	1.76	1,16	0.92	3.68	1,94	) ()	- •
464	450612	AA682452	1,00	0.57	1.75	1,15	1.61	4.08	1.37	0.55	- ,
465	884539	AA629804	0.19	0.11	1.75	1.41	1,45	3.15	1.71	505	~ •
89	504791	AA152347	20.33	11.60	1.75	96'0	1.89	0.27	2.07	۳. رو در رو	- •
467	281442	N47902	₩.86	2.77	1.75	1.78	1.18	1.01	3.28	7.52	
894	325001	W49583	1.56	0.89	1.75	1.28	1.74	4.25	0.81	8.0	- •
469	489594	AA099523	2.16	1.23	1.75	3.54	0.90	1.41	1.65	9 (	- •
470	417777	W88737	1.97	1.12	1.75	4.98	1.15	0.58	0.80	27.	- •
471	277266	N34395	2.46	1.41	1.75	1.52	1.38	3.06	1.93	0.85	- •
472	814341	AA459123	<b>2</b> .	1.06	1.75	2.31	1.29	0.32	0.72	80.4 10.4	- •
473	276871	N39426	1.00	0.57	1.75	3.28	0.79	2.59	1.10	76.0	- ,
474	713129	AA283007	1.69	0.97	1.74	2.26	<del>-</del> 2	0.48	1.14	3.80	- ,
475	206288	H58702	1.59	0.91	1.74	0.78	<u></u>	9.00	0.32	0.30	
476	703577	AA278757	53	0.91	1.74	3.59	2.66	0.74	0.92	OR: 0	
£.	757244	AA426025	14.29	8.20	1.74	1.16	2.01	3.61	0.76	1.1	- •
478	646657	AA205413	0.12	0.07	1.74	1.33	1.89	3.01	1.38	01.1	- ,
479	809719	AA455497	1.58	0.90	1.74	3.53	0.89	0.89	2.12	1.26	
480	1601852	AA989210	2.60	1.50	1.74	1.45	1.29	1.07	1.83	5 5 7	- •
481	284166	N20482	1.05	0.61	1.73	1.00	2.01	4.19	0.50	98.0	- •
482	235909	H52247	0.98	0.57	1.73	1.27	1.15	3.01	Z.	1.70	- ,
483	809374	AA456569	2.00	1.16	1.73	1.55	1.06	3.80	1,40	0.85	- 、
3	753907	AA479351	2.46	1.42	57.7	1.13	2.12	3.99	0.71	0.71	<b>-</b> ·
485	755299	AA496359	6	1.12	Į.	0.98	2.29	96.0	3.61	0.81	
485	A78411	AA670358	30	1.33	12	1.85	0.98	1.37	3.32	1.14	-
787	500035	AA169154	9	0.05	173	0.66	1.68	1.38	3,49	1.43	-
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	1.19	1.24	3.23	3.05	4 02	8,6	2 2 2		(e) +	3. 0	0.00	76.0	3.30	0.89	1.30	1.10	1.06	¥.	99.0	1,31	187	137	 76. C	777	0.0	07.1	3.20	2.48	76.7	05.0	7.35 7.05	4. 6.	77.	B	5.00	67.	8.00	3 6	- P. C	2.40	1.4.	78.0	3.73	2.00	0.42	3.42	2.25		
	1.30	1.59	1.86	2.38	3 5	2. 4	0.5	0.20	, c	8 8 1	0.77	1.40	1.01	1.61	3.23	0.38	0.71	1,55	0.91	1.40	233	3 -	<u> </u>		9:	1.36	1.25	1.09	1.03	1,58	8 80 1	1.70	7.43	 	/c.r	n 6	9 C	57.7	9 K	1.37	D8.	1,26	1.72	<del>-</del> .16	5,42	1.32	1.91		
	3.64	9.0	600	27.	÷ 6	0.60	3.62	On: 0	3.	1.19	1.35	4.28	1.15	1,48	1.35	0.50	1.26	3.25	3.79	9.65	3 6	9 9	20.5	0.41	1.27	1.43	0.21	0.48	3.19	3.01	1.05	3.08	3.76	6.90	0.70	2,03	<b>4</b>	0.44	5.10	4.0	1.45	3.38	0.44	3.41	1.07	0.93	0.58	•	•
	1.27	1.00	3	, c	5 C	0.93 6.0	0.82	2.84	5.02	1.12	2.35	1.06	1.80	1.32	1.46	5.09	1.52	1.17	38	9 6	8 6	0.58	16.0	1.23	1.36	0.71	1,20	1.12	1,23	1.16	0.85	1.7	1.24	0.85	<b>자</b>	0.74	0.82	0.71	0.69	0.83	0.60	1.79	1.71	0.82	0.73	4.58	54.0	ř	
	1.21	30.6	9 6	90.	0.66	1.67	1.84	4.03	0.71	1.27	3.09	0.84	1.27	3.25	61.1	1.46	3.97	1 20	7 4	0	4.26	3.03	1.82	3.67	4.10	3.79	2.63	3.32	1.46	14.	0.67	1.42	0.68	0.70	1.13	0.80	28.	1.33	0.72	3.26	3,11	£;	0.76	0.95	2 6		- c	را بار ا	12
	12	÷	21.	1.72	1.72	1.72	1.72	1.7.	1.71	1.7.1	1.7	1.71	17.				2, 7	· •	0.5	5.	1.70	1.70	1.70	1.70	1.70	1.70	1.70	1.70	1.70	1.69	1.69	1.69	1.69	1.69	1.68	1.68	1.68	1.68	1.67	1.67	1.67	1.67	1.67	1.87	167		) o .	20.	
	237	£.5.	6.0	0.30	0.37	1,30	2.28	3.16	9.	1.89	0.42	0.60	28.0	\$ <b>\$</b>		2,5	, . , .	0.0	27.7	0.42	9. 28.	19.38	4.95	3.30	1.89	99.	15.91	1.40	0.59	1.61	1.02	<b>2</b> 9.	0.37	0.30	29.16	1.18	2.85.	2.50	3.02	4.30	17.98	0.82	1.26	. a	6.5	3 6	9.0	2.08	
	80	9 6	8.	0.52	<u>8</u>	2.24	3.94	5.42	0.07	323	27.0	5 5	3 5	? ?	;	3,5	5.5	77.0	90.0	0.72	1.43	32.92	8.41	5.61	3.21	2.82	27.00	2.38	00.7	2.72	1.73	2.77	0.63	0.50	49.12	1.89	4.78	4.20	5.05	7.21	30.10	137	1 .	: 5	3 5	2 :	<del></del>	3.46	
	44201437	A 8 50 670 5	AA386/80	AA447514	R51617	N25344	N40951	R31413	AA156873	AA476240	W39609	AA454562	AA285053	747625	2011	0.04570	61701000	071000	K0/184	AA/068/0	AA504845	AA418627	AA782292	AA042990	AA609731	AA400317	AA488986	R01515	AA678087	AA405800	R06716	AA496348	W70343	R42736	AA505141.	R25403	R39446	AA777555	AA196486	H28091	AA181179	R53943	W86282	A A 0 1 0 2 3 5	M79634	4423031	AA084517	AA416767	
	700E03	50500	860868	782576	39442	265888	277185	135065	502527	77.1323	322786	800503	143477	71317	71517	Deale	07/795	153061	125829	431462	825742	731290	855236	486591	1031903	743353	824723	123028	430720	742115	126474	755274	345849	31366	825822	133179	23819	448417	628338	162310	624271	40027	44 5854	70001	303081	35/788	246600	731311	
	907	90	483	480	491	492	493	494	495	408	497	107		מ קר קר	3 3	ទ្ធ ខ្ល	706	3	Š.	505	<b>5</b> 6	207	508	55	3 5	2 2		1 1	3 2	, v	55.6	517	518	550	520	521	522	523	524	525	526	527	/70	9 6	670	25	33	22	

Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

	. ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~ ~	****	
4.29 1.15 1.25 1.25 1.66 3.52 1.65 3.17 1.08 3.52 3.52 1.08	0.55 0.51 3.66 3.98 2.08 1.33 0.83 1.51	0.89 0.95 0.70 0.70 0.65 0.65 1.32 1.32 1.44	0.99 1.29 1.02 1.02 1.76 0.98 0.98
2.35 2.35 2.35 2.35 2.35 2.35 2.35 2.35	0.49 1.33 0.46 1.66 0.20 0.65 1.00 1.19	2.04 2.04 1.00 1.00 1.01 1.47 1.18 1.02 1.03	1.10 1.40 1.40 1.66 1.11 1.29 1.24
0.24 2.04 1.48 4.55 0.70 0.70 0.32 2.59	6.16 1.20 1.20 0.94 0.81 1.35 0.73 0.73 1.10 1.10	3.24 1.36 1.30 3.27 3.29 3.29 3.29 3.35 5.35 5.35 7.80	2.6.6.6.6.6.6.6.6.6.6.6.6.6.6.6.6.6.6.6
0.70 1.02 2.97 2.97 1.35 1.35 0.94 1.07 1.07	0.58 1.44 1.05 1.05 2.94 1.26 0.70 0.70 0.95	0.80 0.80 0.80 0.96 0.96 1.77 1.76 1.25 3.33 3.33	0.48 1.17 1.00 0.97 1.61 1.43 1.24 0.95
1.78 0.60 3.33 1.15 0.67 0.67 1.12 1.30 1.20	3.08 3.14 3.14 3.03 5.00 1.00 3.03 3.03 5.60 5.60 5.60 5.60 5.60 5.60 5.60 5.60	3,59 1,23 1,10 1,16 1,68 1,20 1,20	4.08 0.93 1.18 1.19 1.92 1.30 1.30 1.31 1.41 1.31
1.66 1.66 1.66 1.66 1.66 1.66 1.66 1.65 1.65	26. 26. 26. 26. 26. 26. 26. 26. 26. 26.	<b>2</b>	26.1 26.1 26.1 26.1 26.1 26.1 16.1 16.1
13.12 2.10 0.50 0.50 0.59 0.58 2.17 1.39 1.42 2.14 0.73	5.04 5.04 5.04 5.04 5.04 6.09 6.09 6.09 6.09 6.09	2.06 3.54 0.48 4.20 0.48 0.48 3.20 0.05 0.05 0.05	4 + + + + + + + + + + + + + + + + + + +
21.82 3.50 0.09 0.67 0.98 0.97 44.42 2.31 3.55 33.63	1.25 1.25 1.25 1.26 1.57 1.06 2.33	3.38 5.84 6.88 6.88 6.03 6.13 6.03 7.20 6.57 6.66 6.66	5.73 7.82 7.82 0.09 0.13 5.45 6.43
159658 R77512 AA459110 AA402760 R38678 R52796 AA872257 N77263 AA439034 AA439034	A459293 AA001841 H19327 AA629117 AA702327 AA489219 R36587 N63727 AA44099	T74023 AA530415 AA599741 R53951 AA488976 N52837 AA480870 AA480669 AA019320 N59808 AA61332	AA788970 AA678975 N50843 AA505082 AA446866 AA703219 AI017240 AA431571 W52355 W47000
76169 145491 814320 727028 23443 147249 245624 490647 763796 841309 54061	810901 427896 50926 1035765 447404 825060 137275 293539 845380	84560 854633 1070062 40136 824720 244391 840990 363055 288959 1048789	1240220 383528 280950 825689 784216 435992 1636251 782446 325526 325111 109316
E	<b>4</b> 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	555 555 557 558 568 568 568 568 568	567 568 568 568 570 571 572 575 575 576

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0.84	Ξ:	1.24	37		25.	400	2.00		1.26	5.23	3.15	કુ. ફુ. !	3.27	0.79 6.69	0.43	وي. وي	24.0	5 C	7:27	1.30	0.63	0.39	0.86	1.23	4.07	0.78	0.79	\$	1.16	1.45	0.84	0.35	55.5	3 5	3 6	0.63	33.	6	3 20	3.5	28.0	20.0	0.00	3.56		
1.23	1.06	80.	5.00 7.00 7.00 7.00 7.00 7.00 7.00 7.00	5 6	0.03	. ·	1.46	4.51	1.30	1.45	2.13	= :	2.23	1.70	0.74	0.30		4	1.28	<del>-</del> 3	7.00	1.19	0.76	1.01	1.36	3.26	1.17	0.97	0.95	1.07	2.24	0,75	0.99	07.C	0.72 70 0		1.98	96.0	2,00		10.6	D 7		t €	9	
 1.05	5		7.7	8 c	60.0	0.53	3.86	0.19	3.05	<u></u>	0.93	0.78	0.30	3.47	3.54	3.18	3.80	3.47	0.44	0.65	3.68	1.10	1.00	0.97	0.22	1.68	4.39	1.05	3.42	3.21	1,03	0.95	0.63	3.91	) o	0 0	5.00	2.00	3.44	0.50	3.55	55.T	1.20	9,79	1.27	
5		 	0.43	1.28	3.17	0.62	1.15	0.47	1.08	0.78	67.0	3.29	<del>1</del> .	0.67	2.31	1.46	2.82	1.65	0.49	1.35	1,75	1.08	1,28	1,24	0.78	1.09	1.13	1.63	1.72	0.75	0.50	4.88	0.85	0.79	S: 5	1.28	88.0	<u> </u>	5.30	0.67	1.12	1.08	0.8 <b>4</b>	0.50	0.87	
08	60.0	3.39	.05	<u>,</u>	2.11	4.05	9.64	1.10	1.29	122	\$	1.44	1,12	1.30	0.90	1,02	0.52	1.16	3.17	3.56	0.82	3.51	197	3.42	1 43	1.05	0.38	3.17	0.59	1.36	3,13	0.87	3.66	0.60	3.78	3.72	1.92	1.13	1.18	1.06	1.05	0.12	3.99	3.52	o.73	14
9	00.	1.60	1.60	1.60	1.60	1.60	1.60	1.60	1.60	160	1.60	.59	1.59	1.59	1.59	1.58	1.58	58	58	58	- F	75.7	5 6	75	7	75.	75	75	52	1.57	1,57	1.56	1.56	1.56	1.56	1.56	<b>-</b> .56	1.56	1.56	1,55	1.55	1.55	1.55	1.55	1.55	
	0.39	0.53	4.55	2.15	0.75	1.74	1.09	7.03	55.0	8,00	4 45		3.47	2	70	0.29	66.0	1.75	137	0.43	2		3 6	, c	0.0		 	. c	8 24 8 24	0.47	1 47	3.03	0.30	15.46	0.55	0.71	0.80	2.75	0.69	4.16	0.27	2.70	6.02	3.76	0.24	
•	0.62	0.86	7.28	3.4	1.20	2.78	1.74	11 24	2	8 0	9.38 7.45	2 gg	3 5	7.54	£ 5	) 4k	25	, F	2 4	: g	) <del>-</del>	<u> </u>	<b>V</b>	χ <b>ς</b>	<b>9</b> 5	15.30		, d	\$ 6 5 0	9.00	25.0	4.72	0.47	24.12	0.85	<del>1</del> .5	1.24	4.29	1.08	8.47	0.43	4.19	<b>3</b> .6	5.83	0.37	
-	H29268	T99011	AA521431	R63106	R37519	AA478967	N21338	AA001449	N2457B	AA44686	T55728	AA187207	AA056232	A A 434092	AA677534	AA021628	R52901	R24506	R26960	H74265	A A 4 4 3 8 3 2	A A C C 2 4 7 2	D38057	150830£	NS1303	A A 100273	D7478	NABA70	A A 4 5 4 6 3 4	AA214031	WSORAS	AA677287	AA705319	N47240	AA282134	AA505051	AA778653	H23202	R12808	AA035171	R38017	AA620715	R44447	AA876225	H96557	
	49842	122684	826173	137940	137254	753982	285103	181074	2018/4	150104	25570	674400	37777	97770	450403	364083	41560	434704	433273	12026	704065	20000	852598	24855	196387	37536	627627	00454	24/000	657305	244005	454774	462856	280459	711918	825654	1049009	52079	25384	471892	24131	1049230	34869	431505	250988	
	578	579	88	188	582	2	3 2	202	n 9	8 3	è	000	8 6		603	707	200		) ) )			8	86	3 3	5 S	<b>3</b> 5	3 3	ŠŠ	3 8	3 8	200	9 9	9 6	611	612	613	614	615	616	817	618	618	620	23	22	

0.037 1.10 0.033 0.034 0.035 0.0 Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample. 1.21 0.095 0.080 0.070 1.24 1.130 0.070 0. 11.12 9.083 9.083 9.141 1.123 1.103 1.128 1. <del>ૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢૢ</del> **ૢૢૢ** 0.80 1.04 4.21 0.48 0.69 0.75 0.03 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.08 0.09 AA598899
R86847
AA489016
A418595
AA499016
H15895
AA499437
R51912
N51335
N52178
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AA159578
T60149
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AA290137
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T74105
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897928 165828 824920 49203 49203 49203 39593 283142 77284 284247 284247 286247 360478 1472743 149742 149742 149743 149743 149742 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149743 149883 149843 149843 149843 149843 149843 149843 149843 149843 149883 149843

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9	4.43 7.43	<u> </u>	7 C	3 5	0.52	1.17	50.	1.65	2	080	979		. 25	6	10	3.17	0.93	101	0.72	76.	<u> </u>	5 6	5 6	0 k	07.5	0.20	6.7		<b>5</b> 8	96.0	0.00 #	. <del>.</del>	08.0	60	3.41	0.87	0.72	- E	6.0	800	3	5 6	0.35	: E	3	
	0.76	0.27	1,75	 	28.0	3.20	97.0	0.43	200	£.07	5.4	S. 5	S . •	3, 4	2 5	89.0	360	123	5.5	6.6	7°.	) ()	4 60	1.28	0.98	C.83	0.81	e./8	0.73	9. 50 8. 50 8. 50	0.0	0.60	0.67	8 6	60.	80 0	23.		28.0	5 5	S 6	£0.7	 	7/1		
	0.50	0.29	3.40	3.49	24.5. C. C. C.	2.33	51.1	2.0	700	3 6	C 60	300	9.5 8.6 8.6	5.0	<b>3</b> 7 7 7	<b>t</b> t			80°-	C1.7	3.16	3.27	0.35	3.63	0.80	0.74	3.88	422	<b>3</b>	0.87	0.72	\$ 6	8 T		- E	3 6	3 5	3.5	5.5	3.01	3.93	3.14	0.89	8.0 8.0	3.87	
	1.01	6.15	1.05	1.25	0.58	<b>3</b>	1.31	3,64	1,45	66.0	1.01	1.45	0.97	C .93	0.73	78.0	9.6	. i		3.02	1.05	1.02	1.49	0.97	9. 28.	1.15	0.79	0.81	1.12	0.88	1.45	1.19	1.05	 		\$ \$	3.35	0.85	0.82	1.28	0.67	0.69	9.	9. 13.	1.08	
	0.77	0.39	0.44	0.55	0,37	1.21	0.63	0.87	3.20	0.20	3.51	0.57	3.68	3.07	0.56	1,13	.5. .5.	5 !	0.77	0.85	0.90	90.1	3.96	0.94	1.47	1.24	1.14	06.0	3.49	3.61	3.33	3.07	3.02	3.96	0.62	60.0	0.74	1,12	0.58	1.39	0.65	69.0	3.01	1.06	0.94	16
	1.49	1.49	1.49	1.49	1.49	1.49	1,49	1,49	1.49	1.49	1.49	1,49	1.48	1.48	1.48	1,48	1.48	1.48	1.48	1.48	1,47	1.47	1,47	1,47	1.47	1.47	1.47	1,46	1.46	1,46	1.46	1.46	1.46	1.46	1,45	1.45	1.45	1,45	1.45	1,45	1.45	1.45	1,45	1.45	1.45	
	3.07	0.78	5	<b>2</b> .	10.74	0. 4	1.16	1,44	0.82	6.83 83.	0.53	0.41	4.56	0. 12.	15.21	0.98	0.34	6.02	0.50	5.35	125	0.80	0.45	1.40	6.52	90'9	0.48	0.79	0.59	0.49	0.70	0.39	0.63	0.15	<b>24</b> .04	5.67	0.39 ·	4.70	92.77	0.55	2.27	<b>5</b> .	1.31	7.46	50.1	
	4.59	117	1.51	2.46	16.03	0.65	5.7	2.15	1,23	10.16	0.78	0.60	6.76	1.09	22.54	1.46	0.50	8.90	0.74	1.99	<b>4</b> .	1.18	0.68	5.06	9.58	7.47	0.67	1.15	0.86	0,72	<del>.</del> 8	0.57	0.92	0.22	¥.	8,23	0.57	6.82	134.60	0.80	3.30	1.52	1.90	10.78	1.51	
	W16832	AA598817	N35888	W87826	AA703161	AA454928	R08121	AA873578	AA131406	R26785	N33443	R41544	AA400457	AA609392	AA702077	R12385	H03436	N69528	AA489729	AA017468	AA962119	AA701900	N58417	AA707400	H08899	N30285	AA644211	AA678203	AA283949	W86214	R2644	H19320	AA405569	AA453085	H84657	T88721	R52526	AA701046	H62839	AA213669	R93591	R66082	AA233070	AA488064	R26206	
	320392	897956	272529	417081	435577	814738	127120	1472152	503617	132636	270750	34007	753162	743481	384224	128165	150897	294503	823663	361291	1600281	435536	248117	451546	44975	257504	845477	430854	700571	415868	133150	50918	772425	789069	219888	109863	38977	397432	208377	683278	187731	140174	666492	840686	133053	1 1 1
	889	3 8	8 6	67.1	672	579	674	675	676	22	878	679	88	681	682	<b>683</b>	<b>684</b>	88	989	687	889	8	99	ě	68	69	8 8	695	98	269	889	669	8	5	ğ	8	ğ	<b>2</b> 65	902	707	208	802	2	E	712	!

Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

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0.41	3.19	0.85	0.78	2.63	1.22	3.17	1.06	0.87	0.98	3.05	4.02	0.46	<del>2</del> .	3.03	1.22	0.29	1,20	3.04	0.24	1,48	76.0	0.81	20.	0.93	1.63	0.58	1.18	3.44	£.05	0.85	3,13	0.46	0.82	0.92	0.82	16:0	ម្ចី រ	D.73	7 5	<b>2</b> 2	1.02	0.82	1.13	1.32		
1.61	2.98	0.91	0.81	0.68	1.07	1.00	0.86	1.16	1.33	0.42	0.89	0.81	0.62	11	1.03	0.47	1,18	0.67	0.24	0.60	880	94	3.22	62.0	1.27	0.58	0.98	0.65	1.12	0.82	1.22	0.83	0.82	1.12	1.28	1.09.	1.25	0.38	1.07	1.19	1.36	0.95	1.14	3.08		
163	0.42	3.03	60.6	13	0.72	, c	99.0	3.27	505	66	65.0	4.45	) i	65.0	69 6	4 80	3.28	53.0	2000	0.53	5.6	7		60. 60.	3.03		30.E	. 44.0		4 15	0.71	3.26	3.38	3,16	1.72	3.16	3.32	0.51	3.14	3.29	3.21	3.59	3.34	0.64		
ç	0.70	5 C		, c	0.70	 	67:	50. 30.	5.7	- 60	5. 6.	200	2 6	o •	73.0	5 .	67.0	4 6.	2	9 6	3.07	g (5)	0.92	1.02	באים	0.92	28.0	<b>8</b> 6	17.1	2.0	0.50	\$ 50 5 7	101	8	3.00	1.01	0.64	1.63	0.79	0.65	0.91	06.0	0.85	0.92	}	•
•	0.47	C .	CG: 1	5.10	3.00	3.26	0.85	3.63	0.60	3.19	1.37	0.65	0.63	3.19	41,1	0.51	0.31	0./4	1.74	5.74	1.40	0.99	3.28	0.72	0.67	3.32	4.06	0.70	£	0.64	0.67	 	5.50	97.0	0.73	98.0	0.56	3.77	0.79	65.0	15.0	22.0	7.0	, K	2 2	<i>/</i> L
	1.44 1.44	4.	4	<u>.</u> 4	4	<u>-</u> .	4.	4	1.43	1,43	1.43	1.43	1.43	1.43	1.43	1.42	1.42	1.42	1.42	1.42	1.42	1.42	1.42	1.42	1.42	1.42	1.42	1.41	1.41	1.41	1,41	1.41	1.4.	- ·	1.4	- ·	14	•	140	0.4	9 9	9	9.4	5.5	04.1	
	3.41	72.70	1.08	156.38	43.22	0.48	0.22	1.28	1.87	0.26	7.16	1.03	0.44	0.83	0.49	29.18	4.77	3.62	7.29	9.12	7.88	1.20	0.79	2.28	10.38	25.35	0.39	2.54	0.92	3.02	2.74	4.75	<del>-</del> .	2.44	7.07	18.21	2.70	) ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ; ;	2.7	3.43	5.49	 •	3.01	33.92	0.65	
	4.93	104.94	1.58	225.26	62.25	0.69	0.32	<del>2</del> .	2.68	0.37	10.24	1.47	0.64	1.18	0.71	41.58	6.80	5.16	10.37	12.96	11.19	1.70	1.10	3.20	14.70	35.80	0.55	3.59	£.7	4.25	3.85	6.70	1.97	3.44	1.50	25.61	3.88	28.28	2.99	<b>?</b> :	14.69	0.20	421	47.45	0.92	
	N67766	AA872383	AA455659	AA683050	R47979	N63286	W90067	T97890	R76505	AA099820	T90980	AA974971	R39066	AA437241	H70047	AA708240	AA625890	AA677457	R25823	N47316	H87106	N75318	H99215	R43485	AA704332	AA668470	R42533	AA707121	AA448599	AA001199	AA284534	173556	AA480906	N51740	H60696	T62040	N58052	081/02	AA432124	40000	NZ4155	AA481052	. H62529	H92216	AA460282	
	291426	. 1472735	814018	971367	153411	290162	418081	121530	143759	489800	112482	1555523	24915	. 757451	239448	392607	744944	460218	36393	280528	252382	298779	261851	32495	383706	853809	30275	452059	785975	362251	713133	82734	814595	281737	208985	82609	247448	126847	781467	451918	269563	814661	208165	221778	785730	
	713	714	715	716	2	718	749	22	2	2	<u> </u>	2	2	28	727	728	2	8	2	23	2	3 2	i k	3 2	3 2	2 2	3 8	9	7	742	743	4	745	<b>3</b>	747	<b>2</b> 8	49	8	75	32	K	75	33	758	157	

				•		•	•		- •	- •	- •	_	- ,	- •			•	• •			•	•			-		-	-	•	· •-	• •		· -	_	-	-	-	-	-	-	-	-	-	-		•	
ć	, e	1,24	1.14	0.78	50	-	1 5	1 E	27.	2.0	9 5	50.7	CO. 4	1.21		2.0	6 5	82.0	1.58	1,10	3.68	0.72	99	27.0	0.87	1.28	1.90	0.70	0.73	9	0.83	86.0	9. 3.	1.28	 4:-	0.86	3.18	1.40	1.18	0.43	0.47	1.09	9.0	0.54	5	<u>:</u>	
ř	0.70	127	0.84	0.80	080	78 C	108	26.0	9 6	- c	67.1	8.75 0	000	0.78	90 <del>-</del>	3 6	74.0	88.0	0.64	130	0.11	1.08	- C	122	8	76.0	0.65	0.88	1.71	0.39	0.70	4,06	1,15	0.93	1.24	0.47	0.89	0.31	1.39	0.30	0.57	0.67	0,60	1.73	66.0	<u>}</u>	
9	0.0	3.27	3.62	3.10	3.47	98	3.47	3.24	25.0	30.0	5 6	87 O	000	3.20	5.50	1 22	3.68	0.90	3.06	3.14	60	163	3 82	3.07	3.26	3.06	0.35	0.82	3.09	1.74	1.13	0.38	3.03	3.35	3.07	0.57	0.53	3.38	3.07	4.54	3.54	3.19	0.91	3.16	3.07		
, u	00.1	0.60	0.74	1.39	0.89	0.75	0.80	101	÷ +	900	6 6	# 9 5 •	9	5	2.0	0.53	0.83	0.73	0.92	0.75	1.31	3.06	28.0	1.03	0.79	0.82	0.86	0.88	0.95	3.17	3.37	0.74	0.77	0.71	0.78	1.27	1.06	26.0	9.64	0.86	1.71	0.86	0.95	72.0	0.83	<u>;</u>	
C 25	3	0.60	0.63	0.89	0.75	0.63	0 49	0.97	3.88	90.0	3 6	) C	n 6	0.62	) t	1 10	0.61	3.81	0.66	0.57	0.85	0.36	0.48	0.80	0.83	0.72	3.05	3.53	0.26	0.50	0.77	9.0	1.20	0.50	0.54	3.60	1.1	0.69	0.46	0.59	0.43	0.71	3.41	0.49	0.78	18	
9.70	2	1.40	1.39	1.39	1.39	1.39	1.39	1.39	1.39		. •		 	0.1.4 as 4		85	138	88.	1.37	1.37	1,37	1.37	1.37	1.37	1.37	1.37	1.36	1.36	1.36	1.36	1.36	1.36	1.36	1.35	1.35	1.35	1.35	1.35	1.35 S.1	<u>수</u> 왕	¥.	<u>۔</u> لا	<b>1.</b> 3	1.34	£.7		
7, 47	5 ;	15.21	19.67	0.09	1.32	34.88	52.40	1.03	0.42	5	80.0	5.45	7 2 0	0.63 4 27	1 29	2.02	0.73	0.52	0.57	102.66	0.10	13.04	1.68	0.70	3.31	13.58	1.57	0.92	7.82	26.33	2.60	1.37	2.32	19.13	24.47	80.0	1.61	0.13	51.24	0.67	1.16	5.76	4.40	69.0	0.37		
8.67	3	21.24	27.27	0.13	<u>~</u> \$	48.51	72.84	£.	0.59	1 40	288	3 2	5 5	1.76	1.79	2.79	5	0.71	0.78	140.80	0.13	17.85	2.29	96.0	4.52	18.54	2.14	1.25	<b>10.62</b>	35.81	3.53	1.86	3.15	25.91	33.12	80.0	2.18	0.17	69.06	0.91	1.56	57.7	200	0.92	0.49		
R78607	036790	H84/39	AA702193	AA054439	AA598775	H81716	H65300	H04757	R17747	AA678361	R56840	R00707	AA682565	N92895	R44837	AA430367	AA430410	R19878	AA704749	H94870	AA776702	N62394	AA682642	AA431772	R70598	AA279172	AA620608	AA443116	AA291972	AA425692	AA447764	AA434403	AA428341	H92215	AA454925	A4433348	2080CN	612001	AA (100.40)	AA83/485	K45118	H79078	AA253442	H92875	W44938		
144932	240037	758817	384.24	381054	836058	219861	208225	152270	25396	431866	138255	123474	431269	309081	33854	769857	769947	34888	451161	230236	1292829	288663	450680	782501	141854	704076	1048696	809488	725364	773204	813661	770879	769673	221778	814744	921000	27.040	262514	29202	1384851	35010	248734	669367	231718	322857		
758	750	e i	8	781	ğ	<b>763</b>	ğ	765	<b>3</b> 6	797	768	769	22	Ē	E	E	7.	217	276	E	138	2	780	781	782	£	\$	<b>8</b>	. <b>788</b>	787	<b>88</b>	789	£ 3	16/	78,	2 2	į	5 g	8 8	<b>≥</b> 6	2 1	82	8	8	<b>20</b>		

Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

<del> +</del>		<del>-</del>	-	_	<b>-</b> ,	<b>,</b> ,		- •				_	_	-	<b>-</b> -	-	<b>-</b> ·	<b>,-</b> ,	<b>-</b> ,	- •			. 🕶	-	-	-	-	<b>.</b>		- 🕶		· <b>-</b> -	_	_	_	-	-	<b>-</b>	-		
0.33	1.05	3.01	1,14	0.65	3.11	0.70	98:0	0.59	9.00	ئ. د د د	0.00	0.83	1.48	0.95	0.91	0.84	1.13	0.55	1.02	1.15	 	S C	0.82	0.98	0.60	0.87	1.16	0.75	1.20	0.00	5 C	1.20	1.15	0.69	0.62	0.85	1.08	0.70	0.68		
0.58	0.62	00.	50,	0.51	1.02	3.43	1.10	0.55	0.82	0.77	57.1 30.0	0.93 20.5	25.0	0.45	1,19	0.85	0,76	1,33	0.73	1.01	0.95	8 8 9 9	0.82		1.18	1.17	1,02	0.68	0.48	99.0	) (0.00	0.32	880	0.41	.63	0.76	0.87	0.44	0.38		
4.83 3.01	% % %	3.30	3.45	3.26	0.86	0.97	0.45	0.51	3.86	0.46	က နှ	3.63	04.0 04.0	0.30 45.00	3.16	3,49	0.23	3.39	0.89	3.01	0.31	3.25	0.62	4 5	79.0	 	00.6	3.29	. 0.21	3.46	3.77	3.32	(C.D.	5.0d 2.0d 2.0d	2.0.4 DE 6	3.50 5.50 5.50 5.50 5.50 5.50 5.50 5.50	90.5	3.30	0.93		
0.57 0.52	3.32	5.6	0.92	100	0.75	0,68	0.91	0.54	0.80	0.70	0.73	0.55 44. (	0.70	0.44	£ 6	0.70	0.73	0.58	0,81	0.69	0.93	0.67	9.64	0.80	1,14	0.0	0.02 0.63	69.0	1.39	980	1.08	0.74	99.0	79. G	7.0	0 ay	9	3 7	3.67	; ;	
0.35 0.46	0.52	0.67	1.04	- C	- C	080	3.14	. 4. 85.	0.42	1.19	0.77	0.71	0.56	3,13	0.73	6.0	2.63 3.68	890	30° E	0.63	3.14	0.54	3.76	0.78	3.02	0.63	4.0	866	3.10	0.51	0.51	0.50	3.13	0.61	0.55	65.0	5.0 6.0	0 0	0.50	• •	91
1.33	1.33	1.33	£	1.32	1.32	25 25 25	70:-	<u> </u>		13.5	1.31	1.31	1.31	1.31	1.30	5. G	0 0	 	 		1.29	1.29	1.29	1.29	1.28	1.28	1.28	97.7	77	1.27	1.27	1.27	1.27	1.27	1.26	1.26	1.26	1.26	07:1	3	
2.54 78 95	1.93	26.74	0.52	4.55	0.40	0.38 0.38	2 6	57.7	2.5		0.50	5.91	11.97	0.34	11.49	3.82	10.08	55.4	21.7	17:7	8 6	9.3	0.93	0.86	1.09	1.15	3.49	35.89	0.63 80.6	0.21	2.88	0.32	9.82	23.43	2.13	1.28	2.16	17.73	0.80	1.72	
3.38	2.57	35.64	0.70	6.02	0.13	0.51	3 3	3.01	5.30	2. c	0.65	7.73	15.65	0.45	14.99	4.99	13.13	5.71	1.46	98.5	2 GB	12.02	1.19	=:	1.40	1.47	4.47	46.13	0.83	2,6	3.67	0.40	12.44	29.64	2.68	1.60	2.72	22.29	<del>-</del> 5	2.15	
AA676920	W3/332 AA497111	H66675	H09172	AA682838	AA758470	AA432134	A444/592	AA417921	AA6/5138	AA401626 AA464979	AA504505	AA976561	AA704908	AA927544	AA485303	AA677337	H69553	163324	A427551	K00283	H11732	AA701328	AA164836	AA778663	H15549	AA481437	AA704370	AA197334	169593	AA485249	N47443	AA774501	AA157813	R66923	AA677300	AA482127	N\$0829	N76088	AA486755	N59532	
480164	322136 823562	211387	46518	450300	431029	781492	782687	767405	1472589	725224	801018	1558655	451363	1552618	840266	454469	212473	80109	770059	123278	70CR97	435428	595148	1049030	49499	756554	383752	645079	83541	0.000	280697	399264	588915	140299	454795	756450	280982	285454	841057	248631	
803	\$ §	808	807	808	808	810	811	812	813	814	0 4 0 4	24	818	819	820	821	822	823	824	825	828	3 6	828	3 2	834	832	833	\$	8 8	3 8	3 8	3 8	8 8	2	842	25	3	845	846	847	

Table 2-7; Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 1 tumor sample.

	gan gan gan gan yan yan gan gan gan gan gan gan 'A	************	
0.76 0.91 0.96 0.60 0.76 3.03	0.67 3.01 4.02 0.00 0.73 0.73 1.16 1.10 0.82 0.82	0.75 0.74 0.83 0.83 1.34 1.04 1.25 0.69	9.54 9.55 9.55 9.55 9.55 9.55 9.55 9.55
0.40 0.61 1.05 0.88 0.89 0.27	0.79 0.79 1.85 0.73 0.73 0.73 0.82	0.67 0.67 0.68 0.45 0.78 0.03	0.65 0.65 0.05 0.05 0.00 0.04 0.45 0.65 0.65
9.34 9.25 9.28 9.40 0.70	3.00 3.00 3.00 3.00 3.00 3.00 3.00 3.00	9.28 9.28 9.38 9.38 9.05 9.05	3.26 9.24 9.29 9.30 9.30 9.31 9.32 9.30
0.000 000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.	0.08 0.96 1.71 1.71 1.107 1.12 0.69 0.74	0.64 0.65 0.65 0.65 0.65 0.65 0.65 0.65 0.65	1,11 0,96 0,73 1,12 1,105 1,105 1,05 0,05 0,05 0,05 0,05 0,
3.84 1.04 0.79 0.56 3.33 3.33	3.17 3.17 3.17 3.14 9.63 9.10 9.53	0.48 0.63 0.54 0.60 0.62 3.12 3.33 3.33	0.23 0.22 0.22 0.22 0.44 0.77 0.77 0.77 0.77
<b>3</b> 2 3 4 <b>3 5</b> 5 5	<b>333333333</b> 33333	<b>22 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2</b>	5 5 5 5 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7 7
4.19 1.47 5.09 2.25 15.13 2.25 18.72	3.79 1.35 1.35 5.17 5.17 0.09 0.05 0.05 1.12 1.12 1.78 1.78 1.78	3.92 0.58 4.94 14.12 31.16 5.54 8.11 2.96 0.95 16.23	3.68 12.63 13.05 2.63 1.20 0.02 0.02 2.14 7.58 6.99 10.12
5.25 1.84 5.37 2.82 18.89 2.31.31	4.71 1.89 1.68 0.04 0.07 3.57 13.72 21.84 5.30	4.80 0.71 6.03 17.18 37.89 6.64 9.69 7.79 7.79	1.44 1.502 1.36 1.36 2.31 2.31 10.45
R34225 AA491261 AA707080 AA213667 H86559 AA451863 R92281	AA454986 AA465194 H54364 AA001444 N32071 AA425298 AA701976 AA701976 AA701976 AA701976 H29500 N29457 R55965 H13438	AA757711 AA703159 H38848 AA504132 N21688 A1015195 R34273 AA704278 AA699719 R16146	AAC1364 H10072 AA478284 AA625979 R84454 T83245 AA057195 AA621384 N49107 AA625666 AA478470 N91767
136026 824270 451696 683274 223180 786308	811914 815090 203350 361943 260168 1404995 773284 435953 460143 52730 270331 40638	396085 435596 190692 825234 266455 1623328 136070 450997 433300 53158	123755 46843 754280 745480 141361 80050 489076 504358 826995 279800 745347 786802
848 850 851 853 853	855 860 861 863 863 864 865 865	858 870 871 872 873 875 876	888 888 888 888 888 888 888 888 889 991

Table 2-8: Good vs. Poor Clinical Outcome: Genes upregulated 3 fold or more in at least 2 tumor samples.

Table 2-8: Good vs Poor Clinical outcome: Genes upregulated 3 fold or more in at least 2 tumor samples

$\cdot$		Gen Bank		- 1	EnidayaGand/Av Fold/mpm27/Av Fold/mpm28/Av Fold(mpm29/Av Fold/mpm30/Av Fold(mpm36/Av	old/mom27/Av	-old(mpm28/Av	Fold(mpm29/Av	old(mpm30/A	Fold(mpm35/Av	# of Tumora
Order		Accession	AveGood	AvePoor	ePoor)	ePoor)	ePoor)	ePoor)	ePoor)	(Look)	200
		Number		;		707	CR.R.	3.51	99'0	7.30	4
-	289679	W05628	2.55	0.50	20.6	0.0	3.61	6.33	4.17	6.26	4
7	811162	AA486471	28.87	6.85	17.4	2 5	2.20	4.22	2.10	21.15	m ,
m	471641	AA034945	11.03	1.08	10.17	51.13	9	0.27	9.85	7.08	כיז
4	281039	N47717	36.67	4.42	8.29	22.87	 	740	7 05	8.80	ო
V)	282089	N48259	27.13	3,45	7,87	22.41	÷ ;	. ·	9.52	15.90	n
•	155072	R71393	3.32	0.45	7.4	8.50	78.	00.0	5 30	10.17	60
۰ ۲	208210	H65286	2.70	0.42	6.48	14.00	1.98	9 6	5	5.96	က
. 60	770068	AA430545	26.72	4.62	5.78	12.05	0.67	35.0	8 Y	12.92	က
, o:	712950	AA282273	38.22	6.94	5.50	8.84 4	2. S	C. 53	, e	3.28	m
5	81331	T60111	4.13	0.82	5,03	11.18	1.30	0.00	88.	5.42	n
: =	287728	N59158	3.49	0.71	4.90	14.39	0.35	0.40	3.5	3.71	n
. 5	41228	R58953	0.05	0.01	4.38	9.07	2.53	, c	: C	3.18	m
<u> </u>	122345	199191	0.03	0.01	4.16	12.56	4.17	6.7.9	5 5	9.31	m
4	186918	H43317	6.27	1.60	3.93	3.28	1.4.	6.7	2.55	5.03	n
Ť.	1049033	AA778575	2.50	0.65	3.87	4.86	2.38	1 B	18.6	6.23	m
9	195365	R88904	1.81	0.50	3.62	ල. වෙර	64.4	0.00	2.28	0.54	n
4	502.4	H16743	1.48	0.43	3.43	5.36		7 22	1.32	1.35	n
18	838611	AA456975	87.41	25.99	3.36	5.71	77.4	1 5	3.24	3.77	က
\$	283196	N45282	5,16	1.55	3.32	7.31	. o. e/	75.0	4.76	3.74	n
2	745525	AA626248	4.16	1.27	3.28	5.86	1.7.1	, c	9	3.11	e
7	159487	H15926	0.12	9. 2	2.92	2.50	2000	) r	52.0	3.97	n
8	1034473	AA779728	76.12	27.06	2.81	2.22	3.21	2 4	200	3.28	m
2	13844	R88272	2.02	0.74	2.75	4.16	8 8	6.0	3.13	3.31	m
7	378488	AA777187	17.01	7.17	2.37	3.23	D. 12	56.9 AA AC	1 49	1.02	7
×	767130	AA424535	4.28	0.37	11.65	1.60	20.07 7.4.7	44.45	3	1.22	7
<b>58</b>	307337	N95226	10.81	0.98	11.29	69.	4	33.58	2	3.17	7
23	502625	AA134576	7.27	0.30	8.08	4. 4	£ 5	15.74	8	0.71	7
28	41869	R66438	1.41	0.28	5.03	 0	3,4	2.93	66.0	7.09	7
58	301735	N908B2	4.10	0.90	4.56			11,13	1,94	2.65	2
ន	136605	R35051	32.22	65.7	95.4 95.4	6.3 6.6	2.34	0.78	6.56	7.23	7
<u>ب</u>	362278	AA001219	5.53	1.41	- S- C-	5 5	6 50	10.09	2.88	1.01	~
32	756708	AA443803	2.58	0.67	3.84	9.00	8C.4	11.96	0.59	0.98	2
ង	265028	N21084	1.22	0.32	3.78	0.0		11 75	69.0	0.31	7
ಸ	1409509	AA868929	2.84	0.75	3.78	0.50 1.00	0000	10 11	5,36	3.00	2
35	365707	AA025434	<b>2</b> .	0.53	3.68	0.58	9.50	20.04	0.93	0.97	7
×	768953	AA424920	3.43	<b>3</b> 6.0	3.66	0.96	4.0	8° €	200	9.41	7
37	357628	W94121	1.80	0.49	3.65	5.62	6. 6.	.00	7	1.38	~
8	35058	R45192	28.75	7.90	3.64	9.19	77.6		!		

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9.44	8.17	4 57	j r	7.13	0.88	3.90	7.68	0.35	3.5	<b>t</b> 6	5.00	3.25	9.18	1.08	8.83	0.75	0 43	1 7	C/:1	<b>.</b>	7.6	2.19	1,24	1.47	4.89	441	75	, c	7/0	57.0	9 6	D 60.00	777	3.62	25.0	8.5	55.6	- 6	وي. دي وي	5 6	. c.	2.50	8.7	ą. 2	71.1	3.25	6.80	3.86	
5.51	70.6		ים פ	2.74	0.99	2.55	1 60	80.0		2.50	2.35	2.42	1.32	990	070		07.0	00.0	1.70	6.81	4.53	1.58	0.82	0.85	4 47	- u	9 5	7.33 7.33	0.46	2.83	0.69	1.56	3.01	2.18	4.00	<u>.</u> 2	3.15	3.69	5.65	1.21	1.21	2.49	1.76	6.54	1.13	1.58	0.28	2.42	
0.73	- 33	70.	2.38	5.00 5.00	10.28	5	9 9	ō. 6	077	8.	0.59	0.91	0.45	8	<u> </u>	<u>.</u> [	0.5	0.74	4.46	9. 8.	1.52	3.16	622		- c	2.5	4 .	<del>.</del> .	7.31	4.22	7.85	8	1.71	S	0.65	6.17	5.37	2.58	60.	5.67	5.03	2.12	<del>1</del> .95	0.61	5.61	1.56	0.22	92.0	٠
0.50		7.	1.91	2.54	3.45	2	60.	4.91	3.72	2.96	1.26	1.48	<u>.</u>	2 4	o F	- 0.0	0.89	4.10	2.99	1.13	1.41	6.21	27.5	2 4	0 6	38.	0.98	1.25	4.28	2.02	0.54	1.22	2.68	1.46	0.37	1.13	0.86	1.20	0.93	3.19	4.91	3.72	3.10	0.53	3.38	2.03	3.78	1.42	
4.47	<b>.</b>	1.63	101	3.72	8	9 6	8.58 1	0.59	8,99	6.07	8.02	7.62	2 8 5	9 6	6.21	3.68	10.04 40.04	0.74	3.39	0.45	0.82	0 0	3 6	3 7	6.94	4.81	0.58	4.87	0.63	3.01	3.32	3.79	3.41	5.42	1.63	3.54	2.51	2.17	0.44	1.63	0.43	0,61	3.24	0.31	0.80	3.47	0.70	3.36	7
	J. 0.	3.51	3.36	3.35		5.63	3.29	3.28	3.26	3.23	3.18	2 3		3.12	3.12	بن =	3.09	3.06	2.86	2.81	284		107	2.77	2.75	2.73	2.72	2.71	2.68	2.67	2.63	2.62	2.62	2.60	2.60	2.59	2.59	2.55	2.50	2.46	2.44	2.44	2.41	2.41	2.41	2.38	2.37	2.36	
į	0.00 0.00	0.02	4.11	2		8.7.O	3.68	2.48	0.25	1.18	222	9 0	00.	0.98	1.30	0.02	0.75	1.36	300	36.00	7.0	2 .	35.15	0.45	0.83	1. 2.	14.62	1.22	1.09	2.61	0.57	6	1.50	1.17	2.46	760	1.63	0.32	15.73	7.91	0.77	1.38	0.48	21.78	0.57	4 A7	. L	5. t	<b>?</b>
	30.23	0.08	13.79		3 6	2.50	12.10	8.05	0.83	3.79	7.05	3 3		3.06	<b>3</b>	90.0	2.32	4.17	1.82	101 20	;	7 50	98.90	<del>1</del> .23	2.29	48.94	39.72	3.31	2.83	6.98	1.50	2 83	3.92	305	88.9	2 44	7	18.0	39.31	19.49	1.87	3.37	1.6	52 42	-	; <del>E</del>		3.55	) 5
	K77848	AA443967	AA633658	A D 4 1 2 4 1 7		5000	H09076	R00276	R61372	AA628113	N53447	A A 775.953	10701	16/128	AA432270	H47076	H02837	AA418748	R45517	NASASO	NEDRIE	010001	AA1808/8	AA700680	H92821	AA487902	T56281	AA878120	T69477	AA496988	N53031	AA150500	T69305	W74533	N74131	H17121	AA702335	R61883	H72722	N34316	AA453526	AA634431	R54590	H53340	N23430	AA479795	H73590	AA021586	
	145740	757200	855575	136.450	204127	140197	46166	123264	37980	1055707	245277	17547	9/8280	66599	782335	193087	151477	788020.	35329	245000	70000	208907	646037	434952	231574	840590	78353	1418621	66902	823655	246430	401751	82905	246587	208447	£30412	2416	42872	232772	277.173	795425	743868	40404	202535	202202	040/07	5000	284022	1
	ဓ္ဗ	<b>4</b>	3		7 !	5	4	45	48	. 4	: 9	9 9	2	ន	2	ដ	S	3	¥2	2 2	2 2	<u>`</u>	2	S S	S S	2	ĸ	:2	. 3	1	: 55	2 5	. 99	2 9	3 5	<b>&gt;</b>	- 5		2	, ite	ç		. a	, ,	٠.	٠ د د	- r	y r	•

4444	2000	<b>, 11 11 11 11</b>	1 11 11 11 11 11	<b>,,,,,,,</b> ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		
1.24 4.33 1.96 4.34	3.45 0.90 1.87	0.6.0 0.0.0 0.8.0 0.8.0 0.8.0	2. 4. 4. 8. 2. 8. 80. 8. 8. 2. 8. 80. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8. 8.	20.0 20.0 20.0 20.0 20.0 20.0 20.0 20.0	3.64 2.6.6.6.6.6.6.6.6.6.7.7.1.6.6.6.6.7.7.7.6.6.6.6	3.47 4.24 3.05 3.05 3.78 3.15 0.61 0.48
3.44 3.80 3.80 3.80	4,10 0.74 4.69	0.24 4.70 3.63 0.51	5.78 6.23 1.06 6.71 5.39	1.60 1.67 2.03 3.46 1.07	2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	3.49 0.58 0.75 0.75 3.37 0.56 0.63
1.42 0.53 1.05	0.30 0.42 0.42	0.11 2.15 0.85	0.43 0.25 0.25 0.97	3.5.0 3.00 5.22 5.43 5.64 5.64 5.64 5.64 5.64 5.64 5.64 5.64	20.2 1.80 1.4.0 1.4.0 1.2.0 1.2.3 1.0.0 1.	0.78 0.78 0.38 0.38 1.87 1.87 3.72 3.89 8.83 8.83
1.57 0.44 1.55	2.10 0.65 1.49 0.83	5.59 1.01 0.82 0.78	0.51 0.52 0.97 3.49	3.05 3.05 0.50 0.93 0.93	3.63 3.63 3.63 3.29 3.29 1.63 1.05	ପ୍ରତ୍ତି ଓ ଓ ଓ ଓ ପ୍ରକ୍ର ଅନ୍ୟ କଥା ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ ଓ
3.94 3.32 3.12	2.84 2.84 2.73 2.4	0.76 0.56 3.06 3.40	0.17 0.32 1.03 3.28	6.0 0.89 0.76 3.99	3.33 9.08 9.09 9.27 6.27 6.28 6.28	1.39 . 1.40 . 1.25 . 1.25 . 0.91 . 0.55 . 0.56 . 0.
2.32 2.31 2.30	2.25 2.25 2.23	2	12222 1222 1242 1444 1444 1444 1444 144	6444444 644444	207 206 207 201 201 201 201 201 201 201 201 201 201	1.96 1.96 1.96 1.89 1.89 1.89 1.80 1.77 1.77 1.76 1.69
0.33 1.04 0.37	<b>6.85</b> 1.26 0.70	5.14 12.67 5.54 1.40 17.15	15.57 13.39 2.98 0.24 1.57	10.50 1.71 3.65 22.41 0.86	0.83 10.94 9.27 3.81 0.01 2.07 1.11	0.85 1.51 2.19 8.31 15.04 13.84 3.24 3.24 0.94
0.77 0.84 0.84	15,54 · 2,85 · 1,57	4.78 28.11 12.27 3.07	34.05 29.18 6.47 0.52	22.38 3.60 7.69 47.09 1.81	1.73 19.09 17.78 0.02 50.71 4.12 2.20	1.67 2.97 4.29 1.5.74 1.5.74 5.85 37.58 37.58 48.33 5.72 1.58
N67039 AA447522 H08862	T67053 N52151 AA456299	A456321 AA663981 AA986345 R76247 H16803	H77766 N57872 AA884926 H50086	R16596 W49563 N72137 AA070226 AA779457 H15877	AA609588 AA111969 AA68897 AA683520 AA459292 N80129 AA03558 R36989	AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
295985 782575 45852	66560 284383 813151	813179 855745 1605539 144747	24716 24717 1468260 179083 770868	129585 324815 291097 530814 1032405 49303	1031701 530185 853998 378813 810899 297392 471682 27098	485854 293282 289337 80948 838500 343887 258300 824340 884606 132702 767449 35804
2 2 2 2	88 89 89	8 2 2 2 3	28 28 28 28 28 28 28 28 28 28 28 28 28 2	<b>5</b> 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	109 109 115 125 125 137	115 116 118 118 122 122 123 128 128

Table 2-9; Good vs. Poor Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Table 2-9: Good vs Poor Clinical Outcome: Genes upregulated 5 fold or more in at least 2 tumor samples.

Orger	IMAGE Clane ID	Gen Bank Accession	Tissue	AveGood	AvePoor	Fold(AveGood/A Fold(mpm28/A Fold(mpm28/Av Fold(mpm38/Av vePoor) vePoor) vePoor) ePoor) ePoor) ePoor)	Fold(mpm27/A vePoor)	Fold(mpm28/A vePoor)	Fold(mpm29/Av •Poor)	Fold(mpm30/Av	Fold(mpm35/Av •Poor)	# OF Tumor Samples
-	281030	MA7747		38.87	4.67	8.29	22.87	1.40	0.27	9.85	7.08	ຄ
. 2	282089	N48250	e in state	27.13	3.45	7.87	22.41	0.41	0.67	7.05	8.80	m
•	155072	R71393		3,32	0.45	7.44	8.50	1.97	1.30	9.52	15.90	m
· •	208210	HR5288	fastia	2.70	0.42	6.48	14.00	1.98	0.98	5.30	10.17	က
w	770068	AA430545	prain.	28.72	4.62	5.78	12.05	. 0.87	1.90	8.30	5.98	n
9	299679	W05828	bone	2.65	0.50	5.07	7.07	6.80	3.51	0.68	7,30	60
-	787130	AA424535	kldney	4.28	0.37	11.65	1.60	25.69	28.44	1.49	1.02	7
••	307337	N95226	spleen	10.81	0.86	11.29	1.65	7,47	44.45	1.65	<del>1</del>	~
•	471641	AA034945	brah	11.03	1.08	10.17	21.19	2.20	4.22	2.10	21.15	2
5	712950	AA282273	37	38.22	4	5.50	8.84	1.04	0.35	4.37	12.82	8
=	41889	R66438	adrenal	<del>1,</del>	0.28	5.03	1.46	5.83	15.74	1.30	0.71	7
- 2	81331	T80111		4.13	0.82	5.03	11.18	1.30	0.80	8.59	3.28	7
<b>5</b>	287728	N59158	•	3.49	0.71	4.90	14.39	0.35	0.46	3:86	6.42	7
<b>*</b>	301735	N90882	esophagus	4.10	0.80	4.56	9.05	2.75	2.93	. 66.0	7.09	8
<b>*</b> 5	811162	AA486471	thyrold	28.87	6.65	4.21	0.70	3.61	6.33	4.17	8.28	~
<b>\$</b>	382278	AA001219	Many	5.53	<del>[</del> ]	3.91	2.64	2.34	0.78	6.56	7.23	2
11	1408509	AA868929	bone	2.84	0.75	3.78	0.36	5.86	11.75	0.59	0.31	~
<b>~</b>	788953	AA424920	•	3.43	280	3.66	0.96	5.41	10.03	0.83	0.97	7
6	357828	W94121	•	1.60	0.49	3.65	5.62	0.55	0.68	2.00	9.41	7
8	195365	R88904	•	1.81	0.60	3.62	6.03	1.45	0.88	3.51	6.23	7
2	145740	R77948	prostate	30.23	8.55	3.53	1.47	0.52	0.73	5.51	27.0	~
8	782335	AA432270	endothellal	4.04	1.30	3.12	6.21	5.84	1.80	99.0	 	~

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

Cell Line: Genes upregulated fold or higher in 3 of 3, 5 fold or higher in Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

Order	IMAGE Clone 1D	Gen Bank Accession Number	MPM1(PrEC)	Ave Cancer Ceil Lines Values	Fold(AveCell line/MPM1)	Fold(MPM2/M F PM1)	Fold(MPM2/M Fold(MPM272/M # of Cancer PM1) PM1) Cell Lines	-old(MPM272Jh PM1)	# of Cancer Cell Lines
			į	ř	591.67	7.33	66.37	1701.31	ო
-	757165	AA443950	0.0	0.0	10:- 50 RE 016	4.57	944.19	9.33	6
~	510631	AA099431	0.0	81.5	00.510	C+ 8	680.66	14.13	ຕ
ю	592707	A A 1 F 0 S 0 S	0.01	1.66	234.30	9 9	57075	10.82	ო
4	220851	H95633	0.0	1.28	196.56	8.10	טיייטיט פט מניי	7.15	က
40	782804	AA448186	0.01	1.81	180.70	8.88	250.00	3 40	67
· «c	838831	AA481770	0.02	2.58	117.17	3.60	344.82	; ;;	י ה
۸ (	1343732	AA725564	0.01	1.12	112.48	13.45	301.13	6 7	o e
- 00	768226	AA424900	0.01	<del>2</del> .	103.67	3.09	302.09	\$ \$	<b>)</b> (*)
<b>o</b>	731048	AA421275	0.0	0.99	99.19	3.74	290.43	3.45	) M
. 0	366039	AA071503	0.03	2.66	86.03	5.01	249.63	25.52	) e7
: ;=	795627	AA459917	0.01	0.51	85.17	3.09	239.88	7.41	) e7
. 5	1161830		0.01	0.71	71.26	3.19	203.18		<b>,</b> (1)
<u>.</u>	731371	-	0.03	2.20	71.17	3.28	205.85	7. 45	) r
. 4	784296	AA447079	0.01	0.66	65.30	45.84	105.50	5 7 AC	) e1
. t	798569	AA460463	0.01	0.35	63.57	16.10	138.5	20.00	) ("
5 \$	348942	W94289	0.01	0.61	61.20	3.73	168.10	97.70	<b>.</b> "
2 4	795858	AA461522	0.03	1.60	60.34	6.29	170.43	5. 5	» «
	131452	B23270	0.45	26.93	59.79	104.24	4.69	6.07	, «
5 5	1558655	_	0.08	4.38	54.32	97.06	17.92	) P. /d	<b>,</b> (
2 5	743016		0.01	0.50	50.47	15.70	123.80	S 5	, L
3 8	39442	R51617	0.10	5.02	49.93	99.52	44.7	9 C	) m
22	33076	R44048	0.80	36.21	45.35	56.38	29.19	5 6	) m
ឧ	626390	AA189106	0.07	3.16	45.18	91.62	20.97	2 E	) e2
24	128083	R09747	0.01	0. 44	44.34	25.03	10.107	20.0	
22	1389018	`	0.02	0.81	43.48	<b>X</b>	120.63	3 5	, er
28	28498		0.01	0.43	43.37	5.62	127.33	3.10	) m
27	757358	•	0.01	0.42	41.72	4.59	109.53	20.5	) e7
28	813633	_	9.0	1.68	41.51	3,55	116.08	76.4	•
;				•	•-				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

365209 / 812170 / 50288   243181   256911   1030613   487661   270626   291057   1455635   42415   108378   108378   108378   42415   768056   42415   768056   42415   768056   42415   768056   42415   768056   420055   430025   430033   430032   44595   84790   430033   44595   884790   430033   44595   430032   44595   430032   44595   430032   44596   44596   44696   4	0	1367900	AA810225	0.01	0.41	41.11	4.81	99.20	19.33	m
AAA56633         0.01         0.30         37.18         16.25         80.79           H17888         0.01         0.32         36.79         12.25         87.45           H9482         0.01         0.36         35.62         5.81         88.28           N30088         0.01         0.25         34.44         5.019         73.72           AA608822         0.01         0.33         33.18         5.81         88.27           AA04336         0.02         0.59         33.57         29.44         9.21           N7215         0.23         0.55         20.19         7.48         9.21           AA783331         0.02         0.40         29.15         8.48         9.21           AA783332         0.01         0.40         29.16         8.49         9.21           AA783339         0.02         0.44         28.33         5.99         65.78           H99813         0.02         0.44         28.33         5.89         65.78           H0224         0.01         0.27         27.21         13.82         36.47           H0234         0.01         0.24         27.34         5.16         55.23	_	366209	AA062985	0.0	0.38	38.18	3.08	106.34	5.1	n
50288         H17888         0.01         0.32         36.79         12.25         87.45           243181         H94482         0.01         0.38         3562         5.81         88.28           256911         M30088         0.01         0.25         34.44         20.19         73.72           1030613         AA60852         0.01         0.23         33.16         5.81         88.28           270626         AA33313         0.02         0.25         32.57         29.44         39.44           270626         AA33322         0.01         0.02         0.25         32.57         29.46         38.30           455825         AA863232         0.01         0.44         29.15         3.44         30.44           109378         A733232         0.01         0.40         29.16         3.10         3.24           109378         A768232         0.01         0.44         28.83         5.99         65.78           255014         H99818         0.02         0.44         28.83         5.96         65.78           26026         AA6986         A469818         0.01         0.27         22.13         7.36         65.78 <td< td=""><td></td><td>812170</td><td>AA456035</td><td>9.0</td><td>0.30</td><td>37.18</td><td>16.25</td><td>80.79</td><td>14.51</td><td>60</td></td<>		812170	AA456035	9.0	0.30	37.18	16.25	80.79	14.51	60
243181         H94482         0.01         0.38         3562         5.81         88.28           256911         N30088         0.01         0.25         34.44         20.19         73.72           1030613         A4608832         0.01         0.29         34.44         20.19         73.72           487861         AAC45438         0.02         0.59         32.57         28.44         38.44           281057         N33331         0.01         0.20         0.59         32.57         28.48         38.24           281057         N33331         0.02         0.40         29.16         34.48         38.44           281057         N33331         0.01         0.41         29.16         34.48         38.44           198382         0.02         0.44         29.16         34.48         38.30         38.34           198383         0.02         0.41         22.13         14.89         38.44           25054         A.41         38.63         38.30         38.30         38.30         38.30           1514         H2524         0.01         0.17         26.14         26.14         39.16           26805         A.41585 <th< td=""><td></td><td>50288</td><td>H17888</td><td>0.01</td><td>0.32</td><td>36.79</td><td>12.25</td><td>87.45</td><td>10.68</td><td>6</td></th<>		50288	H17888	0.01	0.32	36.79	12.25	87.45	10.68	6
256911         N30088         0.01         0.25         34.44         20.19         73.72           487861         A4008522         0.01         0.33         33.18         5.81         8.72           487861         A4045436         0.02         0.59         32.57         29.44         38.44           270626         N33331         0.01         0.32         31.52         8.48         9.21           270626         N33331         0.02         0.40         29.16         34.48         38.44           270626         N32331         0.01         0.40         29.16         34.48         38.44           1455025         A405222         0.01         0.41         29.15         7.73         38.44           109378         177729         0.02         0.44         29.13         7.73         38.47           139180         N57         17.34         27.31         18.07         21.58           25004         H99813         0.01         0.17         26.13         35.03           26505         A441803         0.01         0.16         26.24         7.36         55.36           26806         A41595         A41595         0.01 <td< td=""><td></td><td>243181</td><td>H94482</td><td>0.01</td><td>0.36</td><td>35.62</td><td>5.91</td><td>88.28</td><td>12.68</td><td>n</td></td<>		243181	H94482	0.01	0.36	35.62	5.91	88.28	12.68	n
4477861         A4608822         0.01         0.33         33.18         5.81         92.72           4477861         A4044536         0.02         0.59         32.57         29.44         39.44           270626         N33331         0.01         0.32         31.52         8.48         9.21           270626         N33331         0.01         0.40         29.16         9.49         9.24           1455825         AA863222         0.01         0.40         29.16         9.10         30.24           10378         177729         0.01         0.41         29.13         7.73         38.30           103780         AA86328         0.02         0.44         29.16         9.48         9.21           263014         H98913         0.02         0.44         29.13         19.07         21.58           263014         H98913         0.02         0.27         27.21         13.86         9.51           263014         H98913         0.02         0.27         27.21         13.86         9.51           26304         H9224         0.01         0.27         27.24         27.86         9.50           46896         AA418030 <td< td=""><td></td><td>256911</td><td>N30098</td><td>0.01</td><td>0.25</td><td>34.44</td><td>20.19</td><td>73.72</td><td>9.41</td><td>m</td></td<>		256911	N30098	0.01	0.25	34.44	20.19	73.72	9.41	m
AAQUESTAS         0.02         0.59         32.57         29.44         39.44           NY3331         0.01         0.32         31.52         6.48         9.21           NY2115         0.23         6.65         29.19         24.48         9.21           AA863232         0.01         0.41         29.15         9.10         30.24           T77729         0.01         0.41         28.83         5.99         65.78           H99813         0.62         0.44         28.83         5.99         65.78           H99813         0.02         0.44         28.83         5.99         65.78           H99818         0.01         0.27         27.21         13.82         36.47           H09818         0.01         0.27         27.21         13.82         36.4           H09818         0.01         0.17         25.69         3.46         65.35           H09818         0.01         0.17         25.96         33.86         30.16           H0224         0.13         25.69         35.86         30.16         30.16           H0255         0.01         0.10         0.27         25.84         5.16         55.23		1030613	AA608832	0.01	0,33	33.18	5.81	82.72	11.01	n
N33331         0.01         0.32         31.52         648         9.21           N72115         0.23         6.65         29.16         24.48         9.21           AA86322         0.01         0.41         29.16         3.10         30.24           T77729         0.01         0.41         29.13         7.73         36.47           AA78032B         0.02         0.44         28.83         5.99         65.78           H99813         0.02         0.44         28.83         5.99         65.78           H99813         0.02         0.44         28.83         5.99         65.78           H99813         0.02         0.44         28.83         5.99         65.78           AA418903         0.01         0.27         27.21         1.382         36.04           H02294         0.02         3.11         26.24         7.36         56.35           AA418903         0.01         0.24         25.84         5.18         56.23           AA42139         0.01         0.24         25.84         5.18         56.23           AA42139         0.01         0.24         25.84         5.18         56.23	G	487861	AA045436	0.02	0.59	32.57	29.44	39.44	28.82	
AA862322         0.01         6.65         29.19         24.48         38.30           AA862322         0.01         0.40         29.16         9.10         30.24           T77729         0.01         0.41         29.13         7.73         36.47           AA769328         0.02         0.44         28.83         5.99         65.78           H99813         0.63         17.34         27.31         18.07         21.59           H99813         0.63         17.34         27.31         18.07         21.59           H99814         0.01         0.27         27.21         13.82         36.47           AA46903         0.01         0.15         27.21         13.82         36.47           AA46903         0.01         0.16         27.21         3.46         5.35           H0224         0.02         1.21         25.96         36.46         55.35           AA46535         0.01         0.24         25.59         4.23         56.23           AA400340         0.01         0.02         25.59         4.23         52.20           AA400340         0.01         0.02         25.59         4.46         54.42		270626	N33331	0.01	0.32	31.52	8.48	9.21	76.87	က
AAA58322         0.01         0.40         29.16         9.10         30.24           T77729         0.01         0.41         29.13         7.73         36.47           AA789328         0.02         0.44         28.83         5.99         65.78           H99813         0.63         17.34         27.31         19.07         21.59           H99813         0.01         0.27         27.21         13.82         36.44           R60381         0.01         0.27         27.21         13.82         36.44           R60381         0.01         0.16         26.24         7.36         55.35           AA418303         0.01         0.17         25.69         36.89         13.28           H02294         0.48         12.53         25.84         5.16         55.35           AA418303         0.01         0.24         25.84         5.18         56.21           AA452139         0.01         0.24         25.59         4.23         62.71           AA528386         0.01         0.17         25.59         4.23         62.71           AA452838         0.01         0.00         24.80         34.18         16.28	<b>~</b>	291057	N72115	0.23	6.85	29.19	24.48	38.30	24.80	c
AA789328         0.01         0.41         29.13         7.73         36.47           AA789328         0.02         0.44         28.83         5.99         65.78           H99813         0.63         17.34         27.31         19.07         21.59           N92764         0.01         0.27         27.21         13.82         36.04           R60361         0.01         0.27         27.21         13.82         36.04           R60361         0.01         0.12         26.71         36.05         36.04           H03284         0.01         0.16         26.24         7.36         35.03           H02294         0.04         12.53         25.93         38.89         33.16           R49555         0.01         0.24         25.84         5.18         52.20           AA45538         0.01         0.17         25.93         4.46         54.64           AA400383         0.01         0.17         25.59         4.23         65.14           AA4003840         0.40         10.00         24.80         36.1         65.64           AA4003840         0.40         10.00         24.80         36.1         10.09 <tr< td=""><td>· ·</td><td>1455835</td><td>AA863292</td><td>0.0</td><td>0.40</td><td>29.16</td><td>9.10</td><td>30.24</td><td>48.14</td><td>6</td></tr<>	· ·	1455835	AA863292	0.0	0.40	29.16	9.10	30.24	48.14	6
AA789328         0.02         0.44         28.83         5.99         65.78           H99813         0.63         17.34         27.31         19.07         21.59           N92764         0.01         0.27         27.21         13.82         36.04           R60961         0.12         3.11         26.71         9.46         65.35           AA418033         0.01         0.16         26.24         7.36         35.03           H03284         0.48         12.53         25.93         38.89         30.16           H03294         0.49         12.53         25.93         38.89         30.16           R49555         0.01         0.24         25.84         5.18         58.23           AA452139         0.01         0.24         25.89         4.23         62.71           R49556         0.01         0.17         25.99         4.23         62.71           AA45088         0.01         0.17         25.99         4.23         62.71           AA45088         0.01         0.17         25.99         3.61         62.70           AA45088         0.01         0.17         24.80         3.43         16.28	_	108378	177729	0.01	0.41	29.13	7,73	36.47	43.19	6
H89813         0.63         17.34         27.31         19.07         21.59           NB2764         0.01         0.27         27.21         13.82         36.04           R60361         0.12         3.11         26.71         9.46         65.35           AA418903         0.01         0.16         26.24         7.36         35.03           H09818         0.05         1.21         25.86         33.86         30.16           H02294         0.48         12.53         25.89         33.86         30.16           R49555         0.01         0.24         25.84         5.18         58.23           AA452139         0.01         0.24         25.89         4.23         62.71           R59556         0.01         0.17         25.89         4.25         62.71           AA23686         0.01         0.17         25.59         4.25         62.71           AA23686         0.01         0.17         25.69         4.46         54.64           AA23686         0.01         0.10         25.24         4.46         55.20           AA23686         0.01         0.00         0.10         2.480         3.41         4.35	_	1391682	AA789328	0.02	0. 44	28.83	5.99	65.78	14.72	n
NB2764         0.01         0.27         27.21         13.82         36.04           R60361         0.12         3.11         26.71         9.46         65.35           AA418903         0.01         0.16         26.24         7.36         55.03           H09818         0.05         1.21         25.96         33.86         30.16           H02294         0.04         12.53         25.93         38.89         13.26           R49555         0.01         0.24         25.84         5.18         58.23           AA452139         0.01         0.24         25.89         4.23         62.71           R59656         0.01         0.17         25.69         4.23         62.71           AA453686         0.01         0.17         25.69         4.46         54.64           AA400383         0.01         0.17         25.09         3.61         60.05           AA4009840         0.04         10.00         24.80         34.16         13.61           AA4009840         0.01         0.00         24.51         46.39         16.26           AA4509838         0.01         0.16         23.75         8.09         59.71      <	•	263014	H99813	0.63	17.34	27.31	19.07	21.59	41.29	m
AA418903         0.12         3.11         26.71         9.46         65.35           AA418903         0.01         0.16         26.24         7.36         35.03           H09818         0.05         1.21         25.96         33.86         30.16           H02294         0.04         12.53         25.89         38.89         13.28           R49555         0.01         0.24         25.84         5.18         58.23           AA452139         0.01         0.24         25.89         4.23         62.71           R59556         0.01         0.24         25.89         4.23         62.71           AA45139         0.01         0.17         25.59         4.23         62.71           AA400840         0.01         0.17         25.59         3.416         54.64           AA400840         0.01         0.17         25.99         34.18         13.61           AA400840         0.04         10.00         24.80         34.18         13.61           AA52838         0.01         0.16         23.75         8.09         59.71           AA628038         0.01         0.16         23.91         4.37         17.55 <t< td=""><td></td><td>303180</td><td>N92764</td><td>0.01</td><td>0.27</td><td>27.21</td><td>13.82</td><td>38.04</td><td>31.77</td><td>*</td></t<>		303180	N92764	0.01	0.27	27.21	13.82	38.04	31.77	*
AA418903         0.01         0.16         26.24         7.36         35.03           H09818         0.05         1.21         25.86         33.86         30.16           H0224         0.43         12.53         25.93         38.89         13.26           H0224         0.46         12.53         25.84         5.18         56.23           AA452139         0.01         0.24         25.89         4.23         62.71           R59556         0.01         0.17         25.59         12.08         52.20           AA4010383         0.01         0.17         25.59         12.08         52.20           AA4009640         0.40         0.17         25.09         3.61         60.05           AA4009640         0.40         0.10         24.80         34.18         13.61           AA4009640         0.40         0.10         24.80         34.18         16.26           AA718675 -         0.39         24.51         46.38         16.26           AA718675 -         0.39         24.29         30.26         10.09           AA623938 -         0.01         0.16         23.97         4.98         30.37           R483057 -		42415	R60981	0.12	3,11	26.71	9.46	65.35	5.30	60
H09818         0.05         1.21         25.86         33.86         30.16           H02294         0.48         12.53         25.83         38.89         13.26           H02294         0.48         12.53         25.89         13.26         13.26           R49555         0.01         0.29         25.89         4.23         62.71           R59556         0.01         0.17         25.59         12.08         52.20           AA010383         0.01         0.25         25.24         4.46         54.64           AA236986         0.01         0.17         25.09         3.61         60.05           AA008470         0.40         10.00         24.80         34.18         13.61           AA478675         0.38         9.30         24.51         46.38         15.65           AA78677         0.39         24.51         46.38         15.65           AA78678         0.01         0.16         23.97         44.37         17.55           R48112         0.04         0.84         23.75         8.09         59.71           NA530328         0.01         0.16         22.37         4.98         30.37           AA62342		768056	AA418903	0.0	0.16	26.24	7.36	35.03	36.33	8
H0224         0.48         12.53         25.93         38.89         13.26           R49555         0.01         0.24         25.84         5.18         58.23           AA452139         0.01         0.24         25.89         4.23         62.71           R59556         0.01         0.17         25.59         12.08         52.20           AA010383         0.01         0.25         25.24         4.46         54.64           AA236966         0.01         0.17         25.09         3.61         60.05           AA008640         0.04         10.00         24.80         34.18         13.61           AA4236986         0.01         0.10         24.51         46.38         16.26           AA4236986         0.01         0.10         24.51         46.38         16.26           AA478675 -         0.38         9.30         24.51         46.38         16.26           AA529838         0.01         0.16         23.97         44.37         17.65           AA6230328         0.01         0.16         22.37         4.98         30.37           AA6230328         0.17         3.74         22.30         3.82         51.95     <		46896	H09818	0.05	1.21	25.96	33.86	30.16	13.87	es
R49555         0.01         0.24         25.84         5.18         58.23           AA452139         0.01         0.29         25.59         4.23         62.71           R59556         0.01         0.17         25.59         12.08         52.20           AA010383         0.01         0.25         25.24         4.46         54.64           AA036960         0.01         0.17         25.09         3.61         60.05           AA009940         0.40         10.00         24.80         34.18         13.61           AA009940         0.40         10.00         24.80         34.18         13.61           AA009940         0.40         10.00         24.80         34.18         13.61           AA009940         0.01         0.24         24.29         30.26         10.09           AA009940         0.01         0.16         23.97         44.37         17.65           AA009940         0.01         0.16         23.01         8.09         59.71           AA009940         0.01         0.16         22.37         4.98         30.37           AA009940         0.01         0.16         22.30         3.82         51.95 <td></td> <td>151184</td> <td>H02294</td> <td>0.48</td> <td>12.53</td> <td>25.93</td> <td>38.89</td> <td>13.26</td> <td>25.64</td> <td>က</td>		151184	H02294	0.48	12.53	25.93	38.89	13.26	25.64	က
AA452139         0.01         0.29         25.59         4.23         62.71           R59556         0.01         0.17         25.59         12.08         52.20           AA010383         0.01         0.25         25.24         4.46         54.64           AA036960         0.01         0.17         25.09         3.61         60.05           AA036940         0.04         10.00         24.80         34.18         13.61           AA708675         0.38         9.30         24.51         46.38         16.26           AA629838         0.01         0.24         24.29         30.26         10.09           AA629838         0.01         0.16         23.97         44.37         17.65           R42112         0.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         22.37         4.98         30.37           AA620328         0.01         0.16         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         38.20         16.96           AA025142         0.16         3.52         22.30         16.96         59.72		38344	R49555	0.01	0.24	25.84	5.18	58.23	14.11	e
R59556         0.01         0.17         25.59         12.08         52.20           AA010383         0.01         0.25         25.24         4.46         54.64           AA236966         0.01         0.17         25.09         3.61         60.05           AA009640         0.40         10.00         24.80         34.18         13.61           AA778675         0.38         9.30         24.51         46.38         16.26           AA629838         0.01         0.24         24.29         30.25         10.09           R39191         0.01         0.16         23.97         44.37         17.55           R42112         0.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         23.01         8.11         34.50           AA630328         0.01         0.16         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         38.20         16.96           AA025142         0.16         3.52         22.24         38.20         16.96           AA025142         0.16         27.44         37.52         23.72           R61	_	787856	AA452139	0.01	0.29	25.59	4.23	62.71	9.84	n
AA010383         0.01         0.25         25.24         4.46         54.64           AA236986         0.01         0.17         25.09         3.61         60.05           AA009840         0.04         10.00         24.80         34.18         13.61           AA778675         0.38         9.30         24.51         46.38         16.26           AA778675         0.39         24.51         46.38         16.26           AA629838         0.01         0.16         23.97         44.37         17.65           R39191         0.01         0.16         23.07         44.37         17.65           R42112         0.04         0.34         23.75         8.09         59.71           N63057         0.01         0.16         22.30         4.98         30.37           AA630328         0.01         3.74         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         39.20         16.96           AA025142         0.16         3.52         22.44         37.52         23.72           R61866         0.05         21.43         4.47         56.01	_	41595	R59556	0.01	0.17	25.59	12.08	52.20	12.48	60
AA236986         0.01         0.17         25.09         3.61         60.05           AA009640         0.40         10.00         24.80         34.18         13.61           AA778675         0.38         9.30         24.51         46.38         16.26           AA629838         0.01         0.24         24.29         30.26         10.09           R39191         0.01         0.16         23.97         44.37         17.55           R42112         0.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         23.01         8.11         34.50           AA630328         0.01         0.16         22.30         3.82         50.37           AA43335         0.17         3.74         22.30         3.82         51.95           AA630328         0.17         3.74         22.30         38.20         16.96           T61938         1.90         40.77         21.44         37.52         23.72           R61866         0.05         21.43         4.47         56.01	4	130255	AA010383	0.01	0.25	25.24	4.46	54.64	16.63	က
AA009840         0.40         10.00         24.80         34.18         13.61           AA778675 *         0.38         9.30         24.51         46.38         16.26           AA629838         0.01         0.24         24.29         30.26         10.09           R39191         0.01         0.16         23.97         44.37         17.65           R42112         0.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         22.30         8.11         34.50           AA630328         0.01         0.16         22.37         4.98         30.37           AA025142         0.17         3.74         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         38.20         16.96           T61938         1.90         40.77         21.44         37.52         23.72           R61866         0.05         21.43         4.47         56.01	æ	187972	AA236986	0.0	0.17	25.09	3.61	60.05	11.62	n
AA778675 .         0.38         9.30         24.51         46.38         16.26           AA629838 .         0.01         0.24         24.29         30.26         10.09           R39191 .         0.01         0.16         23.97         44.37         17.65           R42112 .         0.04 .         0.84 .         23.75 .         8.09 .         59.71           N63057 .         0.01 .         0.16 .         22.30 .         8.11 .         34.50           AA630328 .         0.01 .         0.16 .         22.37 .         4.98 .         30.37           AA025142 .         0.17 .         3.74 .         22.30 .         3.82 .         51.95           T61838 .         1.50 .         40.77 .         21.44 .         37.52 .         23.72           R61866 .         0.05 .         0.97 .         21.43 .         4.47 .         56.01	*	30005	AA009840	0.40	10.00	24.80	34.18	13.61	. 26.60	B
AA629838         0.01         0.24         24.29         30.26         10.09           R39191         0.01         0.16         23.97         44.37         17.55           R42112         0.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         23.01         8.11         34.50           AA630328         0.01         0.16         22.30         4.98         30.37           AA025142         0.16         3.74         22.30         3.82         51.95           T61838         1.50         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01	×	349033 ,	AA778675	0.38	9.30	24.51	46.38	16.26	10.88	n
R39191         Q.01         0.16         23.97         44.37         17.65           R42112         Q.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         23.01         8.11         34.50           AA630328         0.01         0.16         22.37         4.98         30.37           AA025142         0.16         3.52         22.24         38.20         16.96           T61938         1.50         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01	80	84790	AA629838	0.01	0.24	24.29	30.26	10.09	32.50	က
R42112         0.04         0.84         23.75         8.09         59.71           N63057         0.01         0.16         23.01         8.11         34.50           AA630328         0.01         0.16         22.37         4.98         30.37           AA025142         0.16         3.52         22.24         38.20         16.96           T61938         1.50         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01		23345	R39191	0.01	0.16	23.97	44.37	17,55	9.98	ო
N63057         0.01         0.16         23.01         8.11         34.50           AA630328         0.01         0.16         22.37         4.98         30.37           R44836         0.17         3.74         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         38.20         16.96           T61938         1.50         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01	14	29585	R42112	Ş	980	23.75	8.09	59.71	3.48	n
AA630328         0.01         0.16         22.37         4.98         30.37           R44836         0.17         3.74         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         38.20         16.96           TF1838         1.90         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01	ನ	84714	N63057	0.0	0.16	23.01	8.11	34.50	26.43	m
R44936         0.17         3.74         22.30         3.82         51.95           AA025142         0.16         3.52         22.24         38.20         16.96           T61938         1.50         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01	œ	55910 /	VA630328	0.01	0.16	22.37	4.98	30.37	31.76	ო
AA025142 0.16 3.52 22.24 38.20 16.96 T61938 1.90 40.77 21.44 37.52 23.72 R61866 0.05 0.97 21.43 4.47 56.01	m	4204	R44936	0.17	3.74	22.30	3.82	51.95	11.14	ო
T61938         1.90         40.77         21.44         37.52         23.72           R61866         0.05         0.97         21.43         4.47         56.01	న	35149 A	A025142	0.16	3.52	22.74	38.20	16.96	11.55	က
R81866 0.05 0.97 21.43 4.47 56.01	^	0006	T61938	1.90	40.77	21.44	37.52	23.72	3.08	ED
	*	3065	R61866	0.05	76.0	21.43	4.47	56.01	3.81	m

lated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prostate

	24.62 9.30 40.27 43.05 20.16 10.71 14.13 10.73 16.88 5.21 34.04 4.08 31.84 32.94 5.91 3.40 5.91 3.54 19.22 5.36 15.00 3.61 5.36 15.00 3.61 5.36 15.00 3.61 5.36 15.00 3.61 5.36 5.36 15.00 3.61 5.36 5.36 5.36 5.36 5.36 5.36 5.36 5.36	7.89 7.89 18.45 33.42 30.57 10.12 24.35 38.83 38.83 28.90 33.82 42.11 11.78 10.67 8.92 36.93 36.93 36.93 37.9 5.09 37.53 16.46 37.53 16.46 39.26	5.07 16.18 5.36 20.94 7.04 4.49 10.20 10.45 4.26 4.26 4.26 4.26 4.26 4.26 4.26 4.26 4.26 4.26 4.26 4.26 4.26 5.84 5.84 5.84 5.84 5.84 5.84 5.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.72 7.84 6.73 7.84 6.73 7.84 6.73 7.84 6.73 7.84 6.74 6.	20.40 20.40 20.40 19.50 18.85 18.24 17.83 17.15 17.15 16.92 16.93 16.94 16.34	0.21 0.25 0.20 0.20 0.19 0.17 0.17 0.17 0.10 0.29 0.29 0.30 0.30 0.35 0.35 0.35 0.35 0.35	0.01 0.01 0.01 0.01 0.01 0.01 0.02 0.04 0.04 0.04 0.04 0.04 0.05 0.04 0.07 0.07 0.03 0.03 0.03 0.03 0.03 0.03	N32904 W90036 H22824 N32542 H2263 R54492 AA883187 AA057195 AA01315 T49365 AA00304 N47012 AA192784 AA609304 N92263 U26769 W93717 H81554 T69270 AA410437
, ~	\$0.45 \$7.05	4.19	10.07	14.24	3.51	0.25	AA883800
· 62	28.45	3.05	24.69	14.41	<u>2</u> .	0.11	476543
· •	15.54	3.07	40.0	14.54	0.14	0.01	A437142
m	, the	0.00		14.61	0.35	0.02	R40434
m	22.27	18 45	3 .	14.04	0.70	0.05	A406061
n	3.17	17.53	2 1 3 c	78.51	0.35	0.02	H97140
ო	15.00	24.61	, 44 44	. 00.61	3,80	0.25	A702335
ы	5.36	5.66	33.67	47.5	0.71	0.05	A401378
en	3.61	39.06	3.35	72.51	0.30	20.0	A045175
က	19.22	22.78	4.69	15.57	) C	3 6	A416545
m	6.49	33.79	6.61	15.63	0.29		2000
<b>n</b> (	6.69	36.93	3.84	15.85	0.40	0.03	4410437
, c	12.25	14.07	21.48	15.94	0.32	0.02	T69270
? (	<b>4</b> 0.5	41.27	3.05	15.96	1.17	0.07	H81554
, c	70.07	15.89	5.81	16.17	11.31	0.70	V93717
, «	) i	30.25	9.38	16.34	0.09	0.0	179466
) e:	5 C C C	5.09	10.47	16.35	0.16	0.0	. 692569
· «	- S.C.	35.95	6.72	16.54	0.21	0.01	126769
• •	2 2 2	8.92	7.84	16.56	0.17	0.01	<b>VB1432</b>
, ez	2 2	10.67	9.15	16.89	0.17	0.01	676998
m	34 82	40.40	y	16.92	0.69	0.0 4	192784
က	4.08	17.68	40°.	17.15	0.63	9.04	47012
n	35	44.78	97.70	17.19	0.86	0.05	509304
ည်	5.21	0.81	25.85	17.25	33.78	1.96	19355
က	16.88	50 0	2, 1, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2, 2,	17.36	. 0.21	0.01	701315
က	10.73	33.87	64.01	17.83	2.95	Q.17	047441
es	14.13	6 80	n ti	18.01	0.35	0.02	057195
ю	10.71	3883	10.20	18.24	0.31	0.02	883187
က	20.16	24.75	555	16.63	0.17	0.0	54492
က	43.05	10.12	, r	19.30	9.19	0.0	54036
က	40.27	11.18	20 %	20.2	2.40	0.12	22563
က	9.30	30.57	20.94	70.07	0.20	500	32542
က	22.42	33.42	5.36	20.40	0.20		£7077
, e	24.02	18.45	18.18	20.42	0.25	0.01	27874
•	8	7.89	5.07		0.21	0.01	/90036
, ,	1.05	20.01	: !	20.90	2	0.91	32904

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

, es	က	ო	e	m	m	m	m	m	m	m	m	m	m	ы	n	ო	က	ო	ы	ю	m	m	es	ю	ю	ø	ಳು	n	က	က	n	m	က	60	
5.25	5.35	23,16	5.12	4.48	20.74	15.62	17,41	22.12	19.45	8.12	18.18	6.86	7.18	11.26	3.21	7:37	24.08	17.09	17.55	23.17	12.88	19.11	16.72	12.09	3.40	20.89	24.11	20.94	<del>7</del> .	12.02	12.92	5.22	7.26	5.14	
31.74	5.86	9.24	29.82	18.97	9.76	11.86	15.99	8.92	12.04	9.42	18.08	17.85	29.16	21.72	28.51	7.81	7.58	13.80	15.47	4.23	18.70	13.18	8.33	19.31	20.61	10.18	5.70	7.88	9.83	15.78	8.10	22.31	13.18	4.33	·
5.44	31.21	5.00	6:39	17.85	10.64	13.47	7.33	9.24	8.23	22.16	3.39	14.72	3.08	5.65	6.31	21,95	5.26	5.65	3.28	8.77	4.34	3.09	10.15	3.60	10.87	3.65	4.58	5.06	17.71	4. 2	11.07	4.43	11.48	21.90	
14.14	14.14	14.13	13.78	13.76	13.71	13.65	13.58	13.43	13.24	13.23	13.22	13.14	13.14	12.88	12.67	12.37	12.30	12.18	12.10	12.06	11.98	11.80	11.73	11.67	11.63	11.61	11.46	11.30	10.87	10.81	- 10.70	10.66	10.64	10.45	4
0:20	1.25	17.58	0.39	0.13	15.63	16.52	0.17	0.13	8.63	1.91	0.26	0.23	1.05	0.17	0.25	3.56	0.12	14.39	0.31	16.58	16.82	3.89	0.31	11.24	0.20	27.64	0.11	0.45	5.68	90.0	0.09	5.93	13.51	12.13	
9.0	0.03	1.24	0.03	0.01	1.1	1.21	0.0	0.01	9.65	0.14	0.02	0.05	90.0	0.01	0.02	0.29	0.01	1.18	0.03	1.38	1.40	0.33	0.03	0.96	0.05	2.38	0.04	0. 20	0.52	0.01	0.01	0.56	1.27	1.16	
AA509189		AA774665			4	AA598974	AA128617	AA016225	W93379	AA885478	AA491227	H19217	AA883181	N34042	AA621355	AA598531	AA644550	AA262211	AA778717	AA476576	AA706901	H09940	AA609002	R44816			AA179161	H67712	AA460299	AA461513	AA453433	AA417920	H58234		
1031446	32109	856289	30850	41406	1035796	898286	502003	359269	415089	1466409	824074	51103	1467166	244050	1055121	897807	845441	686172	1049079	785368	451649	46451	1030798	33839	357138	366971	609228	210820	795746	795840	788209	767403	204483	814769	
66	100	101	102	103	\$	50	\$	107	108	<u>\$</u>	110	111	112	113	114	115	116	117	118	119	120	121	22	123	124	125	126	127	128	129	130	131	132	53	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

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2 of 3 or 10 fold or higher in at least 1
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745019 725308 898073 745490 34134 391987 378452 431242 590500 590500 590500 590500 430375 489213 489213 489213 489213 489213 489213 489213 489213 781047 781047 781047 781047 781047 781047 7811884 897956 141726 324901 78217 624627 491184	×	430928	AA678295	0.74	7.64	10.34	20.21	5.77	50.6	· ·
725300         AA291556         0.02         0.16         10.15         3.39         4.07           808073         AA698787         0.04         0.37         10.00         3.74         20.48           34134         AA628787         0.11         1.05         9.95         1.271         3.07           34134         AL0268         0.11         1.05         9.95         21.35         3.47           39187         A00036         0.01         0.37         9.79         3.40         8.94           37842         AA041197         0.05         0.37         9.79         3.40         8.94           3712A         AA62533         1.28         1.26         9.74         3.45         8.94           4312A         AA76534         0.01         0.10         9.62         3.83         6.58           59070         AA7157         0.02         0.17         9.64         4.99         9.81           4315A         AA7574         0.02         0.17         9.64         4.99         9.81           4315A         AA165A2         0.01         0.10         9.62         3.83         6.58           4315A         AA716A2         0.01 <th< th=""><th><u>ro</u></th><th>745019</th><th>AA626028</th><th>0.01</th><th>0.10</th><th>10.24</th><th>3.65</th><th>22.27</th><th>4.82</th><th>m</th></th<>	<u>ro</u>	745019	AA626028	0.01	0.10	10.24	3.65	22.27	4.82	m
898073         AAGSBF87         O.O.         O.37         10.00         3.74         20.48           745490         AAGSB789         0.11         1.05         9.95         11.71         3.07           34134         R44734         0.51         5.07         9.95         11.21         3.07           381887         AAOGSB78         0.11         1.05         9.95         21.35         3.47           381887         AAOGSB53         0.06         0.37         9.78         5.69         8.56           431742         AAAGSB233         1.28         1.28         9.74         3.40         8.94           431742         AAAGSB233         1.28         1.28         9.74         3.49         8.86           59072         AAAGSB23         1.28         1.26         9.74         4.89         9.81           43174         AAAGSB24         0.01         0.10         9.62         4.89         9.81           430039         NA710564         0.22         0.01         9.62         4.89         9.81           44605         AA701662         0.02         0.16         9.46         4.89         8.81           430079         0.15         1.06 <td>9</td> <td>725308</td> <th>AA291556</th> <td>0.02</td> <td>0.18</td> <td>10.15</td> <td>3.39</td> <td>4.07</td> <td>22.97</td> <td>က</td>	9	725308	AA291556	0.02	0.18	10.15	3.39	4.07	22.97	က
3454         AA625979         0.11         1.05         9.95         12.71         3.07           34134         R44734         0.51         5.07         9.92         21.35         3.47           381887         Alo02656         0.06         0.87         9.92         21.35         3.47           381887         Alo02656         0.06         0.67         9.78         5.68         8.56           37424.2         AR62133         1.26         12.26         9.74         3.45         18.90           580500         Ar415242         0.01         0.10         9.62         3.83         6.58           580720         AA715342         0.02         0.10         9.62         3.83         6.58           433567         AA716524         0.02         0.16         9.52         1.104         10.28           280375         AA716534         0.02         0.16         9.52         1.104         10.28           280375         AA716534         0.02         0.16         9.52         1.104         10.28           28037         AA71654         0.02         0.16         9.42         5.45         10.10           28030         N9777         0.1	: <u> </u>	898073	AA598787	9.0	0.37	10.00	3.74	20.48	5.78	က
34134         R44734         0,51         5,07         9,92         21,35         3,47           391967         Alocoses         0,08         0,57         9,79         3,40         9,94           317452         Alocoses         0,08         0,57         9,79         3,40         9,94           43742         Alocoses         0,08         0,32         9,78         3,40         9,84           431742         Alocoses         0,10         0,17         9,64         4,99         9,81           590500         Alcoses         0,01         0,10         9,62         3,83         6,58           233567         Alcoses         0,01         0,10         9,62         3,83         6,58           233567         Alcoses         0,01         0,10         9,62         3,83         6,58           23356         Alcoses         0,11         0,10         9,48         5,53         11,40           240039         NG778         0,15         1,40         9,48         5,53         11,42           240039         NA70284         1,12         10,61         9,44         5,48         11,44           24656         Accoses         0,05	_	745490	AA625979	0.11	1.05	9.95	12.71	3.07	14.06	m
391987         Altotoses         0.06         0.57         9.79         3.40         8.94           376452         AAAd1197         0.03         0.32         9.78         5.69         8.56           431242         AAAG82533         1.26         1.276         9.74         3.45         18.90           590500         AA15761         0.02         0.17         9.64         4.89         8.56           590500         AA16724         0.02         0.16         9.62         3.83         6.58           230502         AA701652         0.02         0.16         9.62         3.83         6.56           230503         N97113         1.12         1.00         9.68         5.53         11.42           240655         AA701662         0.22         2.04         9.46         6.47         11.42           240655         AA70264         0.22         2.04         9.46         6.47         11.40           240655         AA70264         0.22         2.04         9.46         6.47         11.40           24565         AA204830         0.76         7.14         9.41         3.46         5.77           24565         AA204830         0.76		34134	R44734	0.51	5.07	9.92	21.35	3.47	4.93	က
376452         AAAGU197         0.03         0.32         978         5.69         8.56           431242         AAGS2533         1.26         12.26         974         3.45         18.90           590500         AA157261         0.02         0.17         9.64         4.89         9.61           590727         AA157261         0.02         0.17         9.64         4.89         9.61           433567         AA71662         0.02         0.16         9.57         11.04         10.28           273546         N3274         3.29         31.33         9.53         1.60         9.67         10.79           273546         N3274         3.29         31.33         9.53         11.04         10.28           273569         N47113         1.12         10.61         9.46         5.53         10.79           280375         AA702644         0.15         1.40         9.46         5.47         10.79           280300         N24042         1.12         10.65         9.44         4.56         17.4           280300         N24042         1.12         10.53         9.44         4.56         17.4           280300         N24042 <td></td> <td>391987</td> <th>A1003636</th> <td>0.08</td> <td>0.57</td> <td>9.79</td> <td>3.40</td> <td>9.94</td> <td>16.03</td> <td>က</td>		391987	A1003636	0.08	0.57	9.79	3.40	9.94	16.03	က
343,742         AA682533         1,26         9.74         3.45         18.90           590200         AA157261         0.02         0.17         9.64         4.99         9.81           590727         AA165342         0.01         0.10         9.62         3.83         6.58           43367         AA70652         0.02         0.16         9.57         11.04         10.28           273546         N3274         3.29         31.33         9.53         11.04         10.28           273546         N3274         3.29         31.33         9.53         11.04         10.28           273546         N3773         0.15         1.40         9.48         5.53         11.42           280375         N470564         0.22         2.04         9.46         6.47         10.79           280376         AA702644         0.22         2.04         9.46         6.47         14.40           280377         AA702644         0.16         7.14         9.41         4.56         10.79           280376         AA702644         0.22         2.04         9.46         6.47         11.40           280377         AA678348         0.16         1		376452	AA041197	0.03	0.32	9.78	5.69	8.56	15.10	က
800200         AA157261         0.02         0.17         9.64         4.89         9.81           800207         AA157262         0.01         0.10         9.62         3.83         6.58           800207         AA165342         0.01         0.10         9.62         3.83         6.58           43367         AA70652         0.02         0.16         9.57         11.04         10.28           273546         N3274         3.29         31.33         9.52         11.38         10.79           280375         N47133         1.12         10.61         9.52         11.38         10.79           280309         N90779         0.15         1.40         9.48         5.53         11.40           280300         N24042         1.12         10.53         9.44         5.48         13.22           430373         AA678348         0.76         7.14         9.41         4.56         13.74           48255         AA04565         0.37         3.43         9.32         3.79         15.51           48485         AA04566         0.37         3.43         9.32         3.79         15.51           48488         AA04566         0.31 <td></td> <td>431242</td> <th>AA682533</th> <td>1.26</td> <td>12.26</td> <td>9.74</td> <td>3.45</td> <td>18.90</td> <td>6.87</td> <td>က</td>		431242	AA682533	1.26	12.26	9.74	3.45	18.90	6.87	က
590727         AA165342         0.01         9.62         3.83         6.58           433567         AA706532         0.02         0.16         9.57         11.04         10.28           27346         N3274         3.29         31.33         9.53         4.57         20.10           280375         N47113         1.12         10.61         9.52         11.38         10.79           280375         N470264         0.22         2.04         9.46         6.47         10.79           280302         N24042         1.12         10.53         9.44         5.48         10.79           280302         N24042         1.12         10.53         9.44         5.48         10.79           480573         AA678348         0.76         7.14         9.41         3.48         5.97           494565         AA20430         0.62         5.85         9.41         4.56         7.97           48557         AA678348         0.76         7.14         9.41         3.48         5.97           48565         AA04566         0.37         3.43         9.32         13.48         5.97           481488         AA705047         0.10         0.19 </td <td></td> <td>590500</td> <th>AA157261</th> <td>0.02</td> <td>0.17</td> <td>9.64</td> <td>4.99</td> <td>9.81</td> <td>14.13</td> <td>B</td>		590500	AA157261	0.02	0.17	9.64	4.99	9.81	14.13	B
433567         AA701652         0.01         0.16         9.57         11.04         10.28           273546         N3374         3.29         31.33         9.53         4.57         20.10           280375         N47113         1.12         10.61         9.52         11.98         10.79           303099         N80779         0.15         1.40         9.48         5.53         11.42           269300         N24042         1.12         10.53         9.44         5.48         10.79           269300         N24042         1.12         10.53         9.44         5.48         11.40           269300         N24042         1.12         10.53         9.44         5.48         11.40           430973         AAD6462         0.76         7.14         9.41         3.48         5.97           489213         AAD6466         0.76         0.19         9.32         3.79         15.51           489213         AAD6466         0.02         0.19         9.22         12.49         10.86           481821         AAD6466         0.01         0.09         9.16         12.49         10.86           48188         AAD65047         0.0		590727	AA156342	0.0	0.10	9.62	3.83	6.58	18.44	<sub>60</sub>
273546         N33774         3.29         31.33         9.53         4.57         20.10           280375         N47113         1.12         10.61         9.52         11.98         10.79           303099         N90779         0.15         1.40         9.48         5.53         11.42           48059         AA702684         0.22         2.04         9.46         6.47         14.40           269300         N24042         1.12         10.53         9.46         5.48         13.22           430973         AA678348         0.76         7.14         9.41         3.48         5.97           4489213         AA678480         0.76         5.85         9.41         4.56         7.97           489213         AA678665         0.37         0.19         9.32         3.79         15.51           489213         AA678665         0.37         0.19         9.32         3.79         15.51           489213         AA64666         0.01         0.97         9.22         12.49         10.86           48188         AA70547         0.00         0.97         9.22         12.49         10.86           48188         AA70547         0.		295127	AA701652	0.02	0.16	9.57	1.04	10.28	7.37	က
20375         N47113         1.12         10.61         9.52         11.98         10.79           303099         N9779         0.15         1.40         9.48         5.53         11.42           448059         NA702684         0.22         2.04         9.46         6.47         14.0           448059         NA702684         0.22         2.04         9.46         6.47         14.0           259300         N24042         1.12         10.53         9.44         5.48         13.22           430973         AA678348         0.76         7.14         9.41         4.56         7.97           45565         AA204800         0.62         5.85         9.41         4.56         7.97           4855         AA70477         0.02         0.19         9.32         3.79         15.44           4854         AA404665         0.37         3.43         9.24         10.06         11.24           481488         AA705047         0.10         0.97         9.22         12.49         10.85           481488         AA705047         0.10         0.97         9.22         12.49         10.85           48148         AA705047         0.16 </td <td></td> <td>273546</td> <th>N33274</th> <td>3.29</td> <td>31.33</td> <td>9.53</td> <td>4.57</td> <td>20.10</td> <td>3.91</td> <td>က</td>		273546	N33274	3.29	31.33	9.53	4.57	20.10	3.91	က
20500         NIGHTAN         0.15         1.40         9.48         5.53         11.42           448059         AA702684         0.22         2.04         9.46         6.47         14.40           448059         AA702684         0.22         2.04         9.46         6.47         14.40           289300         NI24042         1.12         10.53         9.44         5.48         13.22           430873         AA678348         0.76         7.14         9.41         4.56         7.97           945565         AA204830         0.62         5.85         9.41         4.56         7.97           345522         VY4377         0.02         0.19         9.32         3.79         15.51           489213         AA045665         0.37         3.43         9.32         17.44         17.24           481488         AA705047         0.10         0.97         9.22         12.49         10.86           481488         AA705047         0.10         0.97         9.15         12.49         10.86           481484         AA705047         0.10         0.97         9.15         14.49         9.15         14.48           4859         A		280375	NA7113	1.12	10.61	9.52	11.98	10.79	5.79	က
448059         AAYOZE684         0.22         2.04         946         6.47         14.40           289300         N24042         1.12         10.53         9.44         5.46         13.22           430973         AA678248         0.76         7.14         9.41         4.56         7.97           45555         AA204830         0.62         5.85         9.41         4.56         7.97           34552         W74377         0.02         0.19         9.32         3.79         15.51           485213         AA045665         0.37         3.43         9.32         3.79         15.51           48733         H04789         0.11         1.03         9.24         10.06         12.44           481488         AA705047         0.10         0.97         9.22         12.49         10.86           481488         AA705047         0.10         0.97         9.15         12.49         11.63           234617         H77727         0.07         0.60         9.16         12.99         8.66           48594         H10192         0.35         3.21         9.15         6.38         4.45           147726         H65584         0.05 <td></td> <td>303099</td> <th>97709N</th> <td>0.15</td> <td>1,40</td> <td>9.48</td> <td>5.53</td> <td>11.42</td> <td>11.50</td> <td>ю</td>		303099	97709N	0.15	1,40	9.48	5.53	11.42	11.50	ю
259300         NZ4042         1.12         10.53         9.44         5.48         13.22           430973         AA678348         0.76         7.14         9.41         3.48         5.97           430973         AA678348         0.76         7.14         9.41         3.48         5.97           445565         AA204830         0.62         5.85         9.41         4.56         7.97           48526         VV74377         0.02         0.19         9.32         3.79         15.51           48524         VA70564         0.11         1.03         9.24         10.06         11.24           461486         AA705047         0.10         0.97         9.22         12.49         10.86           781047         AA446462         1.08         9.89         9.17         4.58         11.53           224617         H7727         0.07         0.60         9.16         12.99         8.66           45694         H10192         0.35         3.21         9.15         6.38         9.45           45694         H10192         0.06         9.16         12.99         8.65           45756         AA558817         0.15         1.44		448059	AA702684	0.22	2.04	9.46	6.47	14.40	7.50	6
430973         AA678348         0.76         7.14         9.41         3.48         5.97           645565         AA204830         0.62         5.85         9.41         4.56         7.97           345552         W74377         0.02         0.19         9.32         3.79         15.51           489213         AA045665         0.37         3.43         9.32         9.36         11.24           489213         AA045665         0.37         3.43         9.32         9.36         11.24           481488         AA056047         0.10         0.97         9.22         12.49         10.86           781047         AA446462         1.08         9.89         9.17         4.58         11.53           781047         AA446462         1.08         9.89         9.17         4.58         11.63           781047         AA446462         1.08         9.89         9.17         4.58         11.63           781047         AA446462         1.08         9.89         9.17         4.58         11.63           7784         H10192         0.35         3.21         9.15         8.45           45634         H46176         0.16         1.44	_	269300	N24042	1.12	10.53	9.44	5.48	13.22	9.62	ი
645565         AA204830         0.62         5.85         9.41         4.56         7.97           346552         VY74377         0.02         0.19         9.32         3.79         15.51           489213         AA045665         0.37         3.43         9.32         9.36         11.24           489213         AA04665         0.11         1.03         9.24         10.06         12.44           481488         AA705047         0.10         0.97         9.22         12.49         10.86           781047         AA446462         1.08         9.89         9.17         4.58         11.63           234617         H77727         0.07         0.60         9.16         12.99         8.66           46594         H10192         0.35         3.21         9.15         6.38         8.45           17784         H46176         0.16         1.44         9.15         6.38         8.45           17784         H46176         0.16         1.40         9.13         13.21         8.45           147726         R6584         0.03         0.26         9.09         3.46         10.73           14772         R65854         0.01		430973	AA678348	0.76	7.14	9.41	3.48	5.97	18.79	e
346552         W74377         0.02         0.19         9.32         3.79         15.51           489213         AA045665         0.37         3.43         9.32         3.79         15.51           489213         AA045665         0.37         3.43         9.32         10.06         11.24           481233         H04789         0.11         1.03         9.24         10.06         12.44           481488         AA705047         0.10         0.97         9.22         12.49         10.86           781047         AA446462         1.08         9.89         9.17         4.58         11.63           234617         H77727         0.07         0.60         9.16         12.99         8.66           46594         H10192         0.35         3.21         9.15         8.17         14.63           47784         H46176         0.16         1.44         9.15         6.38         8.45           487956         AA598817         0.15         1.40         9.13         13.21         8.45           487956         AA598817         0.15         1.40         9.13         13.21         8.45           48717         T50699         0.01 </td <td>_</td> <td>645565</td> <th>AA204830</th> <td>0.62</td> <td>5.85</td> <td>9.41</td> <td>4.56</td> <td>7.97</td> <td>15.70</td> <td>60</td>	_	645565	AA204830	0.62	5.85	9.41	4.56	7.97	15.70	60
489213         AAA045665         0.37         3.43         9.32         9.36         11.24           43733         H04789         0.11         1.03         9.24         10.06         12.44           481488         AAA705047         0.10         0.97         9.24         10.06         12.44           781047         AA446462         1.06         9.89         9.17         4.58         11.63           234617         H7777         0.07         0.60         9.16         12.49         10.86           46584         H10192         0.35         3.21         9.15         8.17         14.83           45584         O.16         1.44         9.15         6.38         8.45           897956         AA58817         0.15         1.40         9.13         13.21         8.45           897956         AA588817         0.15         1.40         9.13         13.21         8.45           897956         AA588817         0.15         1.40         9.13         13.21         8.47           897956         AA58881         0.01         0.05         9.09         3.46         10.73           7817         T50699         0.01         0.05		346552	W74377	0.02	0.19	9.32	3.79	15.51	8.67	က
43733         H04789         0.11         1.03         9.24         10.06         12.44           461488         AA705047         0.10         0.97         9.22         12.49         10.86           781047         AA446462         1.08         9.89         9.17         4.58         11.63           234617         H77727         0.07         0.60         9.16         12.99         8.66           46694         H10192         0.35         3.21         9.15         8.17         14.83           17784         H46176         0.16         1.44         9.15         6.17         14.83           877956         AA598817         0.16         1.44         9.15         6.38         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.45           897956         AA59881         0.01         0.05         8.92         7.38         12.47           7817         T50699         0.01         0.05         8.62         9.14         7.21           49184         AA137072         1.49		489213	AA045665	0.37	3.43	9.32	9.36	11.24	7.35	n
461488         AA705047         0.10         0.97         9.22         12.49         10.86           781047         AA446462         1.06         9.89         9.17         4.58         11.63           234617         H77727         0.07         0.60         9.16         12.99         8.66           46594         H10192         0.35         3.21         9.15         8.17         14.83           17784         H46176         0.16         1.44         9.15         6.38         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.47           141726         R69584         0.01         0.16         9.13         13.21         8.47           141726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.01         8.92         7.38         12.47           7817         T50699         0.01         0.05         8.82         9.14         7.21           49184         AA137072         1.49         12.90         8.67         6.69         13.76           242952         H95638         5.65		43733	H04789	0.11	1.03	9.24	10.06	12.44	5.24	ო
781047         AA446462         1.08         9.89         9.17         4.58         11.63           234617         H77727         0.07         0.60         9.16         12.99         8.66           46594         H10192         0.35         3.21         9.15         8.17         14.83           17784         H46176         0.16         1.44         9.15         6.38         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.47           141726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.11         8.92         7.38         12.47           78217         750699         0.01         0.05         8.82         9.14         7.21           49184         AA137072         1.49         12.90         8.72         6.90         13.76           242952         H95838         5.65         48.92         8.65         6.69         13.76           242952         H95838         0.45         3.90         8.62         5.02         10.99           5         5         48.92         3.		461488	AA705047	0.10	0.97	9.22	12.49	10.86	4.33	n
234617         H77727         0.07         0.60         9.16         12.99         8.66           46694         H10192         0.35         3.21         9.15         8.17         14.83           177884         H46176         0.16         1.44         9.15         6.38         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.45           147726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.01         8.92         7.38         12.47           78217         T50699         0.01         0.05         8.82         9.14         7.21           624627         AA187351         0.56         4.90         8.72         6.69         13.76           491184         AA137072         1.49         12.90         8.67         6.69         13.76           242952         H95638         5.65         48.92         8.65         5.02         10.99           826273         AA520999         0.45         3.90         8.62         5.02         10.99           5         5         5         5<		781047	AA46462	1.08	9.89	9.17	4.58	11.63	11.30	က
46694         H10192         0.35         3.21         9.15         8.17         14.83           177884         H46176         0.16         1.44         9.15         6.38         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.47           147726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.01         8.92         7.38         12.47           78217         T50699         0.01         0.05         8.82         9.14         7.21           491184         AA137072         1.49         12.90         8.67         6.69         13.78           242952         H95638         5.65         48.92         8.66         3.86         6.65           242952         H95638         5.65         48.92         8.65         5.02         10.99           826273         AA520999         0.45         3.90         8.62         5.02         10.99           5         5         5         5         6.69         10.99         10.99		234617	H77727	0.07	0.60	9.16	12.99	8.66	5.84	m
177884         H46176         0.16         1.44         9.15         5.38         8.45           897956         AA598817         0.15         1.40         9.13         13.21         8.47           147726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.01         8.92         7.38         12.47           78217         T50699         0.01         0.05         8.82         9.14         7.21           624627         AA187351         0.56         4.90         8.72         6.69         13.76           491184         AA137072         1.49         12.90         8.67         6.69         13.76           242952         H95638         5.65         48.92         8.66         3.86         6.65           826273         AA520999         0.45         3.90         8.62         5.02         10.99           5         5         5         5         7.09         10.99		46694	H10192	0.35	3.21	9.15	8.17	14.83	4.46	က
897956         AA598817         0.15         1.40         9.13         13.21         8.47           141726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.11         8.92         7.38         12.47           78217         T50699         0.01         0.05         8.82         9.14         7.21           624627         AA18751         0.56         4.90         8.72         6.69         9.17           491184         AA137072         1.49         12.90         8.67         6.69         13.78           242952         H95638         5.65         48.92         8.66         3.86         6.65           826273         AA520999         0.45         3.90         8.62         5.02         10.99           5         5         5         5         6.69         10.99		177884	H46176	0.16	4.4	9.15	5.38	8.45	13.62	m
14726         R69584         0.03         0.26         9.09         3.46         10.73           324901         W49672         0.01         0.11         8.92         7.38         12.47           78217         T50699         0.01         0.05         8.82         9.14         7.21           624627         AA187351         0.56         4.90         8.72         6.30         9.17           491184         AA137072         1.49         12.90         8.67         6.69         13.76           242952         H95838         5.65         48.92         8.66         3.86         6.65           826273         AA520899         0.45         3.90         8.62         5.02         10.99           5         5         5         5         70.29         10.99		897956	AA598817	0.15	1.40	9.13	13.21	8.47	5.71	n
324901         W49672         0.01         0.11         8.92         7.38         12.47           78217         T50699         0.01         0.05         8.82         9.14         7.21           624627         AA187351         0.56         4.90         8.72         6.69         9.17           491184         AA137072         1.49         12.90         8.67         6.69         13.78           242952         H95638         5.65         48.92         8.66         3.86         6.65           826273         AA520999         0.45         3.90         8.62         5.02         10.99           5         5         5         5         7         7         7	_	141726	R69584	0.03	0.26	60.6	3.46	10.73	13.08	6
78217         T50699         0.01         0.05         8.82         9.14         7.21           624627         AA187351         0.56         4.90         8.72         6.30         9.17           491184         AA137072         1.49         12.90         8.67         6.69         13.78           242952         H95638         5.65         48.92         8.66         3.86         6.65           826273         AA520999         0.45         3.90         8.62         5.02         10.99           5         5         5         5         5         5         5		324901	W49672	0.01	0.11	8.92	7.38	12.47	6.90	ო
624627 AA187551 0.56 4.90 8.72 6.30 9.17 491184 AA137072 1.49 12.90 8.67 6.69 13.78 242952 H95638 5.65 48.92 8.66 3.86 6.65 826273 AA520999 0.45 3.90 8.62 5.02 10.99		78217	T50699	0.01	0.05	8.82	9.14	7.21	10.11	n
491184 AA137072 1.49 12.90 8.67 6.69 13.78 242952 H95838 5.65 48.92 8.66 3.86 6.65 826273 AA520999 0.45 3.90 8.62 5.02 10.99 5		624627	AA187351	0.56	4.90	8.72	6.30	9.17	11.70	n
242952 H95638 5.65 48.92 8.66 3.86 6.65 826273 AA520999 0.45 3.90 8.62 5.02 10.99 5		491184	AA137072	1.49	12.90	8.67	69.9	13.78	5.53	က
826273 AA520999 0.45 3.90 8.62 5.02 10.99 5	_	242952	H95638	5.65	48.92	8.66	3.83	6.65	15.48	n
		826273	AA520999	0.45	3.90	8.62	5.02	10.99	9.84	67)
						ĸ				

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

m	0	6	n	n	<b>ئ</b>	6	က	n	က	ဗ	<b>6</b>	က	n	n	က	ဗ	<b>6</b> 0	6	6	6	6	8	6	60	8	6	<b>6</b>	ဗ		e			<b></b>	•	
																												•	••			••	••	• •	
3.90	6.37	14.40	6.83	4.37	8.51	7.71	14.93	15.23	15.35	13.11	4.58	5.12	6,49	10.12	4.48	10.00	11.39	10.84	13.28	11.71	9.05	7.97	11.50	7.15	8.23	5.28	4.93	12.46	12.18	7.87	9.46	14.01	4.48	10.54	
14.21	13.73	6.65	8.33	16.37	11.10	13.64	5.10	3.98	4.12	7.53	15.49	6.62	11.97	9.24	7.24	8.93	4.17	7.66	3.58	8.00	5.22	7.09	5.10	4.96	10.88	4.41	6.94	4.68	6.84	6.65	6,44	4.60	13.99	7.03	
2 of 3 or 10 fold of nigner in at least 1 or 3 carroar our inter-	4.95	3.89	9.78	4.16	5.27	3.04	4.35	5.06	4.69	3.42	3.96	11.86	4.86	3.94	11.54	4.23	7.55	4.60	6.23	3.35	8.56	7.77	6.04	10.53	3.47	12.64	10,35	5.03	3.00	7.30	6.01	3.23	3.25	4.11	
iner in az ieas. 8.51	35	8.37	8.31	8.30	8.30	8.13	8.13	8.09	8.05	8.02	8.01	7.87	7.7.7	7.77	7.75	7.72	7,70	7.70	7.70	7.68	7.61	7.61	7.55	7.55	7.53	7.44	7.41	7.39	7.34	7.31	7.30	7.28	7.24	7.23	ဖ
10 fold of nig		2 2	 	90.0	7.83	0.25	21.34	4.0	61.74	0.28	5.63	35	2.89	09'0	5.81	0.35	24.59	13.42	1.68	261.61	5.69	5.53	0.15	8	5.64	0.20	21.88	23.90	57.27	3.90	0.12	3.80	0.41	5.80	
2 of 3 or	3 5	20.0	3 5	2 6	3	5 6	5 5	1.29	7.67	0.03	0.70	20'0	0.37	0.08	0.75	0.05	3.49	1.74	0.22	20.00	0.75	0.73	0.02	0.01	0.75	0.03	2,95	3.23	7.80	0.53	0.02	0.52	90.0	0.80	
7	116261	AA485265	A4455/80	K60019	AA43646	WD6246	W30210	HK4752	T65907	AA705825	H20524	4445038	H29897	AA625764	A4431741	A A SO 6257	T96718	T55607	W15305	A A R 7 R 4 E D	P96941	AAGUSSGB	N30372	AAD70331	AA434403	T61122	AA701455	N93924	AA488367	H29227	AA477400	N70714	H16833	T62072	
	180	840150	CORDS .	34434	272202	20000	4044B	203275	80399	1292182	K2741	813271	190325	745393	782259	1505919	121251	73596	322461	882510	200402	950690	260035	530820	770879	83666	435076	309288	843163	52704	740620	294273	50787	85670	
;	169	170	<u> </u>	77	3 ;		175	ÊÊ	178	2 2	2 4	3 4	5	- <del>-</del>	3 2	ž ž	3 %	187	, <del>4</del>	3 5	3 5	5 5	5 5	100	3 5	2 5	40,	197	198	8 8	5	}	5 5	<u> </u>	Ì

Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prostate

e Can	cer Cell 1	ines vs. Nor	mal Prosta 2 of 3 or 1(	oc Epiniena O fold or high	Cancer Cell Lines vs. Normal Prostatic Epintelial Cell Lines vs. Normal Prostatic Epintelial Cell Cell Cell Cell Cell Cell Cell Ce	mai Prostatic Epititelia Cell Lines Common 1 of 3 cancer cell lines.	cell lines.		
			•	6	7.22	4.39	11.90	5.38	က
25	251936	H97488	4.68	33,83	4 5	3.04	9.60	8.99	ы
202	796114	AA460952	9. 2.	6.75	7.50	5 45 5 45	7.63	8.78	ო
206	853368	AA663310	2.51	18.03	61.7	. 4 . 6	6.35	10.69	m
207	809567	AA456611	0.0	0.10	6.7	11.72	5.87	3.65	٣
208	283173	N45236	5.03	14.78	0.7	- 1:11 - 2:2	7.83	7.98	က
209	126650	R06944	0.43	3.05	7.05	0,50 88.4	95 6	6.50	m
310	143365	R74203	0.01	90.0	7.03	60.	707	9.20	m
; ;	1637343	AI015359	14.33	100.32	7.00	4./4	. u	11.35	rò
	415908	W86196	90.0	0.44	96.9	3.01	76.0	3.14	ĸ
7 9	7//07	D-SEOAR	90.0	0.39	6.93	6.40	17.20		er:
2 2	284507	0505N	0.53	3.69	6.92	6,49	4.41	, r	. ~
<u> </u>	246696	767677	3.93	27.15	6.92	13.48	5.5g	9 9	
212 213	070040	10171	<b>4</b>	7.46	6.92	4.76	10.31	80.6	
216	814225	AA465236	9		6.85	10.00	5.36	5.19	7
217	433576	AA701645	0.0	2.0	283	3.47	4.92	12.11	ю
218	246659	NS7722	0.37	7.3 <del>8</del>	20.5	000	5.03	12.23	က
219	131316	R23055	1.19	8.14	29.9	27.0	10.24	5.38	က
220	681906	AA256231	0.01	0.08	6.79	6.7	7 78	6.30	ы
, ,	214604	H71230	0.05	0.34	6.79	9.29	, i	4 64	63
3 1	46091	H08595	1.32	8.93	6.79	6.37	<b>5</b> 1	40.97	67
3 8	350110	A & 0.100.85	4.09	27.70	6.77	5.57	3.78		. «
23	2000	20001044	2	OF C	6.72	7.84	4.68	c9./	n (
224	51083	H17139	<b>5</b> ;		6.79	8.02	8.01	4.14	n
22	1031885	AA609723	Ž,	5.0	2 6	0.45	5.02	5.60	m
226	134719	R28287	0.34	2.07	6,09	9, 16 1, 17	99.6	6.95	ы
227	814246	AA465593	9.85	65.62	6.60		4 39	8.44	n
228	950445	AA599092	2.82	18.79	6.65	2 ,	1.65	7.62	ო
300	813149	AA456695	0.70	4.67	6.65	7.85		g 12	ю
3 5	204214	H59203	0.68	4.52	6.64 42	8.09	- 1	1 W Y	67
3 3	4488607	A A B B 46.22	0.01	0.07	6.59	7.52	4.		, ("
5	1400041	A A 207813	5.43	35.52	6.5 42.0	3.80	3.32	12.51	. r
757	C3367	201600	900	0.42	6.54	10.89	5.69	5.03	· "
3	7000	10001	1 12	7.31	6.53	5.27	9.69	6.63	) (
ž	51450	140071	1 7	2,64	6.52	4.93	8.97	2.67	o (
235	281733	N48075	- ·	10 C	, <sub>4</sub>	4.19	4.30	11.00	က
238	50383	H17869	3.43	67.77	3. 6	4.26	5.16	10.01	က
237	160233	H21943	1.30	œ. \$	9.48	2 6	8 32	7.97	ы
238	487297	AA045508	4,13	7.26	6,45	3.07	;		
}					7				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

က	<b>6</b>		6	က	ന	က	ຕ		n	က	ო	ы	ო	ຕ	ო	es	ო	က	က	ო	n	ო	೯	භ	ຕີ	m	ო	m	ო	m	က	ີຕ	ო	m	
9.48	9.25	4.22	4.86	6.63	3.15	11.67	4.91	8.69	10.52	4.60	4.01	4.07	8.79	9.96	5.79	4.83	4.48	5.08	7.86	7.19	5.21	5.37	5.40	5.52	7.53	3.97	9.64	9.01	3,11	7.60	9.58	8.71	9.15	7.58	
5.78	4.97	6.40	3.55	5.78	12.06	3.05	10.41	6.93	5.05	10.49	3.15	10.33	5.59	3.86	5.67	5.90	4.21	5.73	6.77	4.55	6.93	8.09	3.03	7.41	3.15	3.97	4.55	4.93	9.49	5.43	4.46	5.22	4.78	5.30	
8.8	2.00	8.52	10.68	6.67	3.73	4.19	3.58	3.26	3.19	3.63	11.55	4.27	4.19	4.63	6.92	7.62	9.63	7.47	3.55	6.43	6.00	4.66	9.62	5.03	7.03	9.69	3.43	3.67	4.97	4.55	3.46	3.50	3.49	4.53	
6.43	6.41	6.38	6.37	6.36	6.32	6.31	6.30	6.29	6.25	6.24	6.24	6.23	6.19	6.17	6.13	6.12	6.11	6.10	90.9	6.08	6.05	6.04	6.02	5.99	5.90	5.88	5.87	5.87	5.86	5.86	5.83	5.81	5.81	5.80	<b>&amp;</b>
1.70	0.82	4.82	0.62	47.65	0.47	2.79	9.0	4.53	2.35	0.15	31.76	40.75	12.17	4.70	7.33	0.14	3.47	16.66	4.93	90.0	29.59	7.89	96.0	0.07	3.18	1.38	0.10	90.0	2.15	8.00	12.98	43.39	14.21	10.85	
0.26	0.13	0.76	0.10	7.49	0.07	0.44	0.0	0.72	0.38	0.02	5.09	6.55	1.97	9.76	1.20	0.02	0.57	2.73	0.81	0.0	4.89	1.31	0.16	0.01	0.54	0.24	0.02	0.01	0.37	1.37	27.2	7.47	2.45	1.87	
AA034115	AA757564	AA431753	W68162	H02231	N50827	R41972	AA447531	N63744	R11605	R38640	H84871	AA460685	AA447662	R68581	R10284	H18017	AA405800	AA425302	H96392	AA102068	AA459318	AA425404	AA620437	R93875	N29918	N81049	AA453714	H09065	AA099523	W94009	H06113	4A454585	R56870	N59136	
430007	395955	782476	343298	151055					130004																					357531	44255	809535 /	41198	287637	٠
239	240	241	242	243	244	245	246	247	248	249	250	251	252	253	254	255	256	257	258	528	260	261	262	263	<b>78</b>	265	366	267	268	569	270	271	212	223	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

		, ;	,,,	5 80	3.92	7.27	6.22
745360	AA625662	9.50 1	12.20	2 4	05.6	5.86	8.09
74738	T57359	2.36	13.63	07.6	3 .	F 13	6.29
147385	H01332	0.03	0.16	5.76	n i	3 6	10.40
884822	AA669341	41.24	237.42	5.76	3.75	3 :	
705740	AAAED310	0.62	3.53	5.72	10.19	3.51	\$ 10 10 10 10 10 10 10 10 10 10 10 10 10 1
	4460777	0.83	4.73	5.69	6.56	5.66	4.86
070400	110012		5.22	5.68	8.14	3.99	4.93
14/828	K81486	4 6	1 8 1 4	5,65	3.60	5,70	7.68
667598	AA228130	3	9 0	, i	<u> </u>	4.10	8.24
884799	AA629849	2.81	15.89	5,65	8 !		277
45882		9.34	1.94	5.63	5.49	5.03	
705064		1,55	8.72	5.63	3.52	4.25	- -
		. Y	25.20	5.62	6.88	4.48	5.48
140662		) ( f		5.61	330	3.47	10.06
399562		5.	8. 5.		7 70	4.88	6,45
220700		96.0	5.38	5.61	n (	5 . c	3.10
132708		0.45	2.53	5.59	5.63		20.3
51485		.0.07	0.40	5.54	4.68	36.0	3.02
2002		4.40	24.34	5.54	8,15	3.92	40.4
<b>t</b> 0000		68.0	5.08	5.49	3.62	6.29	6.56
0.00		3 6	08.0	5.45	3.31	9.67	3.38
204140		5 4	(¥	5.45	4,98	3.50	7.85
32299		00'-	70:0	77 4	3.64	5.09	7.59
361069		0.08	U.S.		663	5.49	4.14
898097		<u>.</u> Ŝ	5.65	7.5	30.7	, r,	5.87
459947		0.30	1.59	5.37	0.4	9 6	7.07
789204		2.16	11.60	5.37	5.05	5.18	13.7
431245		1.90	10.19	5.37	3.91	5.22	0.0
460114	A A 676840	0.10	0.55	5.34	3.39	7.08	5.55
		0.18	0.94	5.33	3.66	6.07	6.25
300 00120		6.73	35.80	5.32	4. 8.	5.23	6.39
301 35115		1 26	6.70	5.32	3.21	3.61	9.14
		} [	13 64	5 30	6.09	5.91	3.90
303 261714		7	t ::	200	6.79	4.32	4.61
26297		0.02	0.10	7.7	) i	3 70	3.67
		0.17	0.91	5.24	8.75	2.5	108
363103		0.69	3.59	5.21	7.86	80.4	9 1
		0.02	0.12	5.20	5.19	5.01	04.6
307 7023		(6.19	97.60	5 17	4.37	3.47	7.68
			2	:			

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

က	es	က	6	n	က	m	က	60	ຕ	ო	ы	က	60	63	es	က	n	n	кò	m	m	ED.	က	က	က	က	က	ო	m	m	က	m	က	m	
6.97	5,45	6.93	4.75	8.15	4.97	4.67	5.63	4.55	4.34	6.25	4.80	5.33	5.26	4.03	6.82	5.61	6.41	5.90	4.80	5.64	6.54	4.37	6.23	4.69	3.60	5.29	3.08	3.05	3.31	6.55	5.01	3.64	3.51	4.93	
5.27	9.00	5.12	4.23	3.76	3.23	7.05	5.83	6.00	4.38	4.66	5.78	6.66	3.76	5.47	4.74	5.83	3.84	5.34	4.57	5.49	4.24	4.27	4.29	3.94	3.35	4.33	7.24	4.15	4.89	3.95	3.86	5.23	7.07	4.94	
3.25	4.99	¥.3	6:39	3.45	7.16	3.62	4.62	4.64	6.45	4.26	4.49	3.05	6.00	5,36	3.28	3,32	4.46	3.38	5.12	3.32	3.64	5.78	3.85	5.62	7.29	4.52	3.82	6.91	5.91	3.56	5.15	5.09	3.32	4.03	
5.16	5.15	5.13	5.12	5.12	5.12	5.11	5.09	5.06	5.06	5.06	5.02	5.02	5.01	4.96	4.95	4.92	4.90	4.87	4.83	4.81	4.81	4.81	4.79	4.75	4.75	4.71	4.71	4.70	4.70	4.69	4.67	4.65	4.63	4.63	9
2.47	5.19	31.82	0.40	0.24	34.01	2.84	0.77	10.81	23,18	8.96	1.02	23.47	4.55	0.75	24.26	11.53	6.70	2.66	15.	0.90	0.81	15.65	1.22	4.83	5.71	0.45	3.94	32.63	3.65	0.19	3.48	1.69	3.36	0.43	
0.48	1.01	6.20	0.08	0.05	6.64	0.56	0.15	2.14	4.58	1.7	0.20	4.68	0.91	0.15	4.90	2.34	1.37	25.0	0.32	0.19	0.17	3.26	0.25	1.02	1.20	0.09	0.84	6.94	0.78	0.04	0.74	0.36	0.73	0.09	
AA135050	AA703000	N86132	R80387	AA663995	AA450265	H18068	AA885851	AA463924	N93715	AA448160	AA683557	AA504272	AA129552	AA626784	N40939	AA443969	AA459632	AA418918	ÅA678135	AA156599	AA776692	AA620553	AA481757	N90523	N50745	AA883127	AA621019	AA448396	N73301	H97000	T66936	AA046321	H23482	AA682405	
490060	447208	278483	147147	855390	789182	50649	1500388	810264	307119	782766	505864	825411	564803	8777787	277163	757206	810974	767994	431895	502296	1292828	951142	810785	306146	283751	1467028	1056200	781341	292068	249949	66406	_	•	461775	
309	310	311	312	313	314	315	316	317	318	318	320	321	322	323	324	325	326	327	328	329	330	33.	332	88	첧	335	336	337	ğ	339	8	ž	342	848	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

ate Ca	ncer cen	itate Cancer Cell Lines vs. Notifiel r 103 au 10 fold or higher in at le	2 of 3 or 1(	o fold or high	er in at least	2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines	cell lines.	, ,	r
Ž	544664	AA074666	4.89	22.62	4.63	3.06	3.22	7.60	າ (
3,5	595109	AA173926	2.40	11.07	4.60	3.28	5.08	5,45	m (
¥ 5	292933	N69491	0.67	3.08	4.60	4.85	3.14	5.80	r) (
3 2	287749	N62245	1.77	8.14	4.60	4.35	3.56	5.89	m
<b>3</b> 3	4E1936	A 8 600767	0.22	0.99	4.58	6.38	4.05	3,31	m
3	40165	MT4066	15.03	68.72	4.57	3.02	5.75	4.95	ю
<b>3</b>	166687	0000	3 5	20	4.56	5.14	4.81	3.75	n
320	18981	153	5 5	, ye	4.56	4.05	3.27	6.37	ю
321	782800	AA448184	5.5 7.0	63,63	55.4	4.56	3.51	5.59	ო
352	296155	W00895	67.1	50.6	) ¥	2.78	4.39	5.44	က
383	282720	N50079	3.70	16.81	<b>d</b>		80 4	534	m
354	588840	AA157787	3.00	13.59	4.53	3.16	9 6	1 18	, «
355	509479	AA056395	4.33	19.58	4.52	3.27	6.13	o ;	<b>o</b> (
386	413074	A A 707794	0.13	0.60	4.52	3.14	3,41	7.01	m
	30700	DESCO	78.0	3.85	4.51	5.43	3.36	4.75	ы
200	06/60	199794	20.00	113.80	4.48	3.35	5.80	4.30	ო
358	1800/8	AAbasuas	0.63	10 51	4.48	5.20	5.13	3.12	ო
328	814765	AA454947	7.35	10.51	o t	) L	4.50	3.19	ю
38	490548	AA126799	1.22	5.43	4.40	2 0	,	3.76	er.
361	795371	AA453273	2.35	10.45	4.45	4.16	5.42	0.75	יו כ
362	358457	W96114	7.33	32.61	4.45	3,43	5.36	<b>6</b> . 0	? (
363	35,8732	W84486	0.67	2.96	4.43	4.85	4.4	4.00	<b>.</b> 0
3	4627623	AID14704	67 0	2.19	4.43	5.61	3.81	3.87	ю
Š	106/063	100000	, c	25	4.40	3.32	4.59	5.27	m
365	245145	W60015	97.7	15.01	4.39	3.03	3.33	6.82	၈
8	942906	AA486430	<u> </u>	j i	-	r.	3.14	4.26	m
367	1475463	AA857804	1.80	7.88	ה ה ה		3.76	6.16	m
368	359135	AA010128	0.47	2.06	0 °	3 5	3 33	5.10	м
369	198104	R94633	0.21	0.93	65.4	r i	9 6	007	er.
370	249562	H84926	2	0.18	4.37	3.06	5.03	, r	) r
371	308468	N95495	2.29	9.99	4.36	5.17	4.31	80.0	<b>3</b> (
377	324313	AA284108	0.02	0.09	4.34	3.11	6.72	3.19	m
373	280342	N49281	4.13	17.91	4.33	3,32	4.21	5.47	ю
	ACID A	P56055	0.15	79.0	4.32	4.06	4.72	4.18	က
	049036	000001	1 28	5.41	4.31	4.10	3.51	5,33	m
3/5	610007	OROROU	3 6	, c	4.31	3.30	6.39	3.24	м
376	418C4	WS0104	70.0	3 ;		76.7	3.24	5.43	ы
37	258761	N30185	1.26	5.44	06.4	7 (7	, ,	69 1	eri
378	25274	R12804	0.33	1.66	4.29	3.40	n / c	9	•
					=				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

3.43 4.67 3.12 4.40 5.97 3.26 5.22 5.18 3.26 3.14 5.01 4.22 3.31 4.71 4.63 5.40 3.23 8.3 5.02 5.93 5.07 3.07 3.86 4 . G 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines. 3.05 3.07 5.07 5.59 8 15. 5.97 63 4.75 3.59 5.98 3.76 3.72 4.62 8 1.98 £.91 4.1 3.58 5.27 3.90 3.90 3.65 3.26 3.45 3.52 3.43 3.30 3.56 3.67 4.66 3.38 3.18 4.96 4.14 4.24 3.66 3.25 3.72 3.33 8 3.08 8 3.72 33 1.22 4.22 4.21 4.20 8 6.19 4.19 4.18 4.18 4.16 4.14 4.13 4.13 4.12 1.1 Ē. 60. 4.08 4.07 €.0 4.07 4.05 ŝ 15.01 .52 0.45 5.38 9.46 5.68 26.51 5.90 21.16 2.02 30.82 2.60 72.78 0.22 6.85 1.14 4.98 3.57 0. 11. 6.42 1.29 2.26 1.36 0.05 1.66 0.49 17.87 0.28 54. 9.64 5.14 7.53 <u>5</u> 2.09 2.11 2.49 AA126259 W88792 **4A975388** AA679569 AA489247 AA025937 AA452968 AA496360 AA838691 AA053129 AA125825 H44448 **4A024637** W47254 AA459909 R59722 N92478 AA644099 AA489000 T59668 R08935 AA678021 N89861 R05693 **VA705142** N64617 R33303 H23985 AA460571 N64508 W86860 H01926 H11792 1556058 453599 1387760 502767 365157 825089 324255 788566 183771 809421 301867 365641 76196 755301 510170 127925 845380 305538 548693 125183 289945 431803 824739 36180 795830 161592 290378 51469 416390 150135 271006 383 385 385 387 387 88 390 393 394 395 396 391 398 399 400 60 5 និ Ŝ

r Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prostate Ca

625764         AA188366         0.91         3.60         3.8           1654974         AI023724         2.04         8.02         3.6           194305         H50556         0.29         1.14         3.9           32870         R43205         0.34         1.32         3.2           786083         AA448676         2.27         8.89         3.5           613126         AA211448         0.43         1.69         3.6           293576         N69283         4.31         16.90         3.6           77243         AA45852         3.06         11.74         3.6           775435         AA45382         2.90         11.14         3.1           84580         AA45382         2.90         11.13         3.1           84580         AA45382         2.90         11.13         3.1           84580         AA45381         0.02         0.04         3.2           75740         AA426341         5.69         0.05         3.3           75740         AA426342         2.41         8.80         3.2           75740         AA426362         0.25         0.05         0.95           75740         AA										
HOLORANA ALIDIZATA         LATA MISCANA ALIDIZATA		į		6	3.60	3.96	4.14	4.14	3.60	က
1932/34 ALIZALIZA		\$ 1	44166300		8 02	3.94	3.65	3.46	4.71	က
19470 F HOURSON         1.44         1.45         4.45         4.15         3.57         3.19         4.45         4.15         3.57         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68         3.68	_	4/24	AIU23/24	<b>.</b> .	4	3.93	4.53	3.70	3.55	<del>د</del>
780813         AAA48676         0.34         1.35         4.14         4.07         3.57           780813         AAA48676         0.32         1.69         3.92         4.47         3.83         3.47           56312         AAA48676         0.27         16.90         3.92         4.47         3.83         3.47           259576         AAA3612         0.02         0.09         3.86         4.33         3.12         4.12           15520         T16520         1.02         0.09         3.86         4.86         3.09         3.09           75435         AAA5144         0.47         3.85         4.86         3.02         3.47         4.12           75426         AAA5142         3.06         11.74         3.85         4.86         3.03         3.09         3.09           75426         AT5139         2.06         11.74         3.85         4.86         3.03         3.05         3.0	-	4305	H50656		; ;	69.6	3.19	4.45	4.15	က
786003         AA448676         2.27         6.89         3.29         4.77         3.83         3.47           203576         Ned21448         4.34         16.90         3.92         4.72         3.75         4.72           203576         Ned21448         4.34         16.90         3.92         3.28         3.75         4.72           203576         AA436152         3.05         11.74         3.85         4.88         3.02         3.67           77244         T50139         2.66         10.26         3.85         4.88         3.02         3.67           7724         T50139         2.66         10.26         3.85         4.88         3.02         3.67           295710         N72597         1.17         4.47         3.82         4.56         3.33         3.58           295710         N72597         0.01         0.04         3.74         3.02         3.58         3.38         3.58	417 32	2870	R43205	<b>\$</b> .0	20.1	202	4 14	4.07	3.57	m
613126         AA211448         0.43         1.68         3.92         4.71         4.72           293576         N6223         4.31         16.90         3.92         3.28         3.75         4.12           293576         N6223         4.31         16.90         3.92         3.28         3.75         4.12           75435         AA436162         3.06         11.74         3.85         6.38         3.09         3.09           75435         AA436162         3.06         11.74         3.85         4.86         3.02         3.09           81307         AA43612         2.66         10.26         3.85         4.86         3.33         3.36           86450         AA63016         0.01         0.04         3.74         3.23         4.33         3.38           3784         AA7788         0.05         0.34         3.68         3.23         3.39         3.58           3784         AA7788         0.05         0.34         3.68         3.23         3.58         3.58           3784         AA7788         0.05         0.34         3.68         3.23         3.58         3.58           3774         AA77888         0.05 <td>418 78</td> <td>6083</td> <td>AA448676</td> <td>2.27</td> <td></td> <td>2 0</td> <td></td> <td>183</td> <td>3.47</td> <td>63</td>	418 78	6083	AA448676	2.27		2 0		183	3.47	63
293576         N69283         4.31         16.90         3.92         3.26         3.72         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12         4.12         3.12	419 61	3126	AA211448	0. 53.	1.69	3.92	j (	3.75	4.72	e
115230         786627         0.02         0.09         3.86         4.33         3.12         4.12           77244         75039         2.66         10.26         3.85         4.81         3.09         3.69           7724         75039         2.66         10.26         3.85         4.81         3.09         3.69           295710         N/72697         1.17         4.47         3.82         4.85         3.33         3.59           295710         N/72697         1.17         4.47         3.82         4.85         3.33         3.58           868030         AA630016         0.01         0.04         3.74         3.23         4.39         3.58           45384         AA77102         0.25         0.95         3.74         3.07         3.46         4.55           455040         AA77102         0.25         0.95         3.74         3.07         3.48         4.65           75740         AA77104         2.86         0.05         3.74         3.24         3.36         3.56           77389         AA62556         0.16         0.26         3.66         3.24         3.36         3.56           77589         AA45372<		3576	N69283	4.31	16.90	3.92	3.20	2 9		۳
76.4355         AA436162         3.6         11.74         3.85         5.38         3.09         3.09           77.244         75.44         75.44         75.44         75.44         3.02         3.67           813075         AA453622         2.66         10.26         3.85         4.85         3.02         3.67           295710         N72697         1.17         4.47         3.82         4.55         3.33         3.58           295710         N72697         1.17         4.47         3.82         4.55         3.33         3.58           295710         N72697         0.01         0.04         3.74         3.23         4.39         3.58           8R4500         AA57082         0.02         0.34         3.88         3.83         4.03         3.58           45540         AA77082         0.02         0.34         3.68         4.08         3.46         4.65           75740         AA477082         0.03         0.34         3.68         4.03         3.46         3.58           75740         AA47702         2.88         0.05         0.34         3.66         3.23         3.24         3.30           77259		5230	T88577	9	0.09	3.86	4.33	3.12	7 .4	> •
77244         78503         Available 1         Available 2         10,26         3.85         4.86         3.02         3.67           77244         75039         2.66         10,26         3.84         4.81         3.36         3.58           28500         AAA3932         2.90         11,13         3.84         4.81         3.36         3.36           28510         AA77092         0.25         0.95         3.74         3.23         4.39         3.58           455842         AA77092         0.25         0.95         3.74         3.07         3.46         4.65           453842         AA77092         0.25         0.95         3.74         3.07         3.46         4.65           453842         AA77092         0.25         0.95         3.74         3.07         3.46         3.51           455840         AA7704         AA7704         AA7704         AA2704         AA7704			10001	5	11.74	3.85	5.38	3.09	3.09	•
17244         T50139         200         11.13         3.84         4.61         3.36         3.36           285710         AA453632         2.80         11.13         3.84         4.61         3.33         3.58           285710         AA453632         2.80         11.13         3.84         4.65         3.33         3.58           285710         AA77092         0.25         0.96         3.74         3.23         4.39         3.59           37784         AA77092         0.25         0.96         3.74         3.07         3.46         4.65           75740         AA42343         2.66         10.51         3.68         3.33         3.19           75740         AA42434         2.69         20.94         3.66         3.24         3.30         3.19           75740         AA42656         0.16         0.59         3.66         3.24         4.33         3.51           75740         AA42656         0.16         0.59         3.65         3.74         3.30         3.26           75740         AA42656         0.16         0.59         3.65         3.74         3.30         3.26           73599         AA42566 <td< td=""><td></td><td>£</td><td>AA436132</td><td>3 5</td><td>90.0</td><td>3.85</td><td>4.86</td><td>3.02</td><td>3.67</td><td>က</td></td<>		£	AA436132	3 5	90.0	3.85	4.86	3.02	3.67	က
813675         AA453632         2.80         11.13         3.02         4.55         3.33         3.58           285710         N72697         1.17         4.47         3.82         4.55         3.33         3.58           285710         N72697         1.17         4.47         3.74         3.25         4.39         3.59           485890         AA650016         0.01         0.04         3.74         3.25         4.39         3.59           453642         AA771092         0.25         0.95         3.74         3.07         3.48         4.65           453642         AA477092         0.05         0.94         3.66         3.24         3.30         4.45           757404         AA42641         5.69         2.094         3.66         3.24         3.30         3.46           757404         AA42642         5.69         2.094         3.66         3.24         3.30         3.45           757404         AA42642         5.69         2.094         3.66         3.24         3.30         3.45           775299         AA42642         5.73         3.66         3.24         4.05         3.28           289658         N79353		7244	T50139	7.00	0.00	50 6	4.81	3.36	3.36	က
295710         N72697         1.17         4.47         5.02         3.74         3.23         4.39         3.59           884590         AA630016         0.01         0.04         3.74         3.23         4.39         3.59           377884         AA50016         0.01         0.04         3.74         3.07         3.46         4.65           45362         AA577082         0.25         0.34         3.68         3.83         4.03         3.19           15863A0         AA677348         2.86         10.51         3.68         3.33         3.46         3.51           7574Q4         AA67541         5.69         20.94         3.68         3.24         3.30         4.45           77589         AA67556         0.16         0.59         3.66         3.24         3.30         4.45           77589         AA62568         0.16         0.59         3.65         4.71         3.21         3.76           487723         AA64562         1.57         5.73         3.66         3.48         4.43         3.28           487723         AA64562         1.57         5.73         3.66         3.74         3.76           27920 <t< td=""><td>424 81</td><td>3675</td><td>AA453832</td><td>5.90</td><td>51.13</td><td></td><td>1 KK</td><td>333</td><td>3.58</td><td>m</td></t<>	424 81	3675	AA453832	5.90	51.13		1 KK	333	3.58	m
884690         AA630016         O.01         O.04         3.74         3.23         4.65           377884         AA777092         0.25         0.95         3.74         3.07         3.48         4.65           4536242         AA77888         0.09         0.34         3.68         3.89         3.49         3.51           1586340         AA47348         2.86         10.51         3.68         3.33         3.85         3.51           1586340         AA426341         5.69         20.94         3.68         3.33         3.85         3.86           757404         AA426341         5.69         20.94         3.68         3.24         4.40         3.86           757404         AA426342         2.41         8.80         3.66         3.24         4.40         3.28           745286         AA63642         2.41         8.80         3.66         3.24         4.40         3.28           77289         AA436372         1.57         5.73         3.66         3.73         3.72         3.72           288658         NY333         0.26         2.51         3.64         4.05         3.72         3.72           17574         AA67572	425 29	5710	N72697	1.17	4.47	2.02	3 8	7 30	3.59	m
377884         AA777082         0.25         0.95         3.74         3.07         3.48         4.03         3.19           45384.2         AA77888         0.09         0.34         3.68         3.83         4.03         3.19           1586340         AA974348         2.86         10.51         3.68         3.33         3.46         3.51           1586340         AA426341         5.69         2.0.94         3.68         3.33         3.85         3.86           745286         AA62558         0.16         0.59         3.66         3.24         3.30         4.45           77239         AA42842         2.41         8.80         3.65         3.48         4.23         3.28           288658         N79353         0.26         0.95         3.65         3.48         4.23         3.28           487723         AA043529         3.81         14.21         3.65         4.71         3.21         3.76           27920         NF7526         0.69         2.51         3.64         4.25         3.75         3.75           48723         AA645379         3.81         14.21         3.64         4.25         3.75         3.75		4690	AA630016	0.01	0.0	3.74	3.23	EC. +	39 7	
453642         AA77588         0.09         0.34         3.68         3.83         4.03         3.19           1506340         AA47588         0.09         0.34         3.68         3.46         3.51           1506340         AA472641         5.69         20.94         3.68         3.33         3.85         3.85           757404         AA426241         5.69         20.94         3.68         3.24         3.30         445           773598         AA628422         2.41         8.80         3.66         3.28         4.40         3.28           773598         AA428422         2.41         8.80         3.66         3.78         4.40         3.28           487723         AA404552         1.57         5.73         3.66         4.71         3.21         3.02           228658         N79353         0.26         0.96         2.51         3.64         4.05         3.12         3.02           288658         N79353         0.26         0.95         3.62         4.71         3.21         3.02           278648         AA4463879         3.81         14.21         3.64         4.05         3.12         3.19           175744		7864	AA777092	0.25	0.95	3.74	3.07	3.48	6. c	o €
1586340         AA974348         2.86         10,51         3.68         4.08         3.46         3.51           757404         AA426341         5.69         20.94         3.66         3.24         3.30         4.45           757404         AA426342         2.41         8.80         3.66         3.24         3.30         4.45           77359         AA426342         2.41         8.80         3.66         3.48         4.23         3.28           487725         AA043552         1.57         5.73         3.66         3.48         4.23         3.28           288658         N79353         0.26         0.95         3.65         4.71         3.21         3.02           289658         N79353         0.26         0.95         3.65         4.71         3.21         3.02           279920         N79526         0.69         2.51         3.64         4.05         3.12         3.76           279920         N79526         0.69         2.51         3.64         4.05         3.12         3.76           176744         AA5330         2.0         5.25         3.67         3.24         3.57         3.62           256260 <t< td=""><td></td><td>35.47</td><td>AA775888</td><td>60:0</td><td>0.34</td><td>3.68</td><td>3.83</td><td>4.03</td><td>2 0</td><td><b>)</b> . c</td></t<>		35.47	AA775888	60:0	0.34	3.68	3.83	4.03	2 0	<b>)</b> . c
757404         AAAZ6341         569         20.94         3.66         3.33         3.85         3.86           745286         AAAZ6341         569         20.94         3.66         3.24         3.30         4.45           77359         AAAZ6742         2.41         8.80         3.66         3.24         3.30         4.45           77359         AAA21352         2.41         8.80         3.66         3.48         4.23         3.28           28863         N79353         0.26         0.95         3.65         4.71         3.21         3.02           288658         N79353         0.26         0.95         3.65         4.71         3.21         3.02           279320         N7526         0.69         2.51         3.64         4.05         3.12         3.76           279320         N7526         0.69         2.51         3.64         4.05         3.12         3.76           279320         N7526         0.69         2.51         3.64         4.05         3.12         3.76           279320         AA45337         AA45337         3.24         3.24         3.27         3.76           280422         AA48344	•	86240	•	2.85	10.51	3.68	4.08	3.46	3.51	n (
745286         AA625658         0.16         0.59         3.66         3.24         3.30         4.45           772599         AA429422         2.41         8.80         3.66         3.28         4.40         3.28           487723         AA429422         2.41         8.80         3.65         4.71         3.21         3.28           28858         N79353         0.26         0.95         3.65         4.71         3.21         3.02           28958         N79353         0.26         0.95         3.64         4.05         3.12         3.02           279920         N5726         0.69         2.51         3.64         4.05         3.12         3.76           17574         R3520         2.0         7.22         3.64         4.05         3.73         3.79           487425         AA453879         3.81         14.21         3.64         4.05         3.73         3.79           487426         AA657129         2.77         10.00         3.61         3.70         3.74         3.70           487427         AA4191461         1.01         3.54         3.56         3.72         3.74         3.72           487277	•			7. 09.	20.94	3.68	3,33	3.85	3.85	<b>.</b>
742586         AMEZDONS         0.10         3.66         3.28         4.40         3.28           773599         AAA28422         2.41         8.80         3.66         3.48         4.23         3.28           487723         AAA43562         1.57         5.73         3.65         4.71         3.21         3.02           288658         NY9353         0.26         0.95         3.65         4.71         3.21         3.02           279920         NS7526         0.69         2.51         3.64         4.05         3.12         3.76           279920         NS7526         0.69         2.51         3.64         4.05         3.12         3.76           15674         R35230         2.00         7.22         3.62         3.26         4.35         3.19           407425         AA646523         2.77         10.00         3.61         3.70         3.74         3.70           487425         AA646523         2.77         10.00         3.61         3.56         3.72         3.71           58947         AA194141         1.01         3.54         3.56         3.73         3.73         3.74         3.72           58045 <t< td=""><td></td><td>5</td><td>150745</td><td>9 4</td><td>0.50</td><td>3.66</td><td>3.24</td><td>3.30</td><td>4.45</td><td>n</td></t<>		5	150745	9 4	0.50	3.66	3.24	3.30	4.45	n
773599         AA428422         2.41         0.80         3.66         3.48         4.23         3.25           487723         AA043552         1.57         5.73         3.66         4.71         3.21         3.02           288658         NY9353         0.26         0.95         3.65         4.71         3.21         3.02           278920         NS7526         0.69         2.51         3.64         4.05         3.12         3.76           13648         AA453879         3.81         14.21         3.64         3.48         4.25         3.76           13648         AA453879         2.00         7.22         3.62         3.72         3.76         4.37           487425         AA046523         2.77         10.00         3.61         3.70         3.44         3.70           487425         AA046523         2.77         10.00         3.61         3.70         3.44         3.70           589479         AA416617         1.40         4.95         3.54         3.05         3.84         3.52           256260         H94617         1.01         3.54         3.52         3.56         3.73           1604586         AA498445		2286	AA625558	2 ;		3.68	3.28	4.40	3.28	m
487723         AA043552         1.57         5.73         5.00         5.74         5.00         5.75         5.00         5.75         5.00         5.75         5.75         5.76         4.71         3.21         3.02         2.86         2.87         4.75         3.21         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.76         3.77         3.76         3.72         3.76         3.77         3.70         3.44         3.70         3.44         3.70         3.74         3.70         3.44         3.72         3.72         3.72	-	88	AA428422	Z.41	0.0	, ,	3.48	4.23	3.25	<b>6</b> 0
28858         N79353         0.26         0.95         3.65         4.71         3.21         3.76           279920         N57526         0.69         2.51         3.64         4.05         3.12         3.76           813548         AA453879         3.81         14.21         3.64         3.48         4.25         3.19           813548         AA453879         3.81         14.21         3.64         3.48         4.25         3.19           487425         AA046523         2.77         10.00         3.61         3.70         3.44         3.70           487425         AA046523         2.77         10.00         3.61         3.70         3.44         3.70           589479         AA167129         4.22         15.14         3.58         3.96         3.72         3.07           256260         H94617         1.40         4.95         3.52         3.59         3.73         3.62           256260         H94617         1.01         3.54         3.52         3.59         3.73         3.24           627273         AA194445         5.65         19.64         3.47         3.47         3.29         3.84         3.52	-	37723	AA043552	1.57	5.73	5.00	; ;	7	3.02	ന
279920         N57526         0.69         2.51         3.64         4.05         3.12         3.12           813548         AA453679         3.81         14.21         3.64         3.48         4.25         3.19           813548         AA453679         3.81         14.21         3.64         3.48         4.25         3.19           13674         RA5530         2.00         7.22         3.62         3.26         3.26         4.37           487425         AA046523         2.77         10.00         3.61         3.70         3.44         3.70           589479         AA167129         4.22         15.14         3.58         3.96         3.62           256260         H94617         1.40         4.95         3.54         3.05         3.86         3.62           256260         H94617         1.01         3.54         3.52         3.59         3.62           256260         H94617         1.01         3.54         3.52         3.59         3.24         3.32           160436         AA986230         0.80         2.80         3.49         3.29         3.84         3.59           275553         R93309         0.19		38658	N79353	0.26	0.95	3.65	V.4	- 25.6	3.7.F	67
813548         AA453879         3.81         14.21         3.64         3.48         4.25           813548         RA453879         3.81         14.21         3.64         3.48         4.25           13674         RA5330         2.00         7.22         3.52         3.26         3.26           487425         AA046523         2.77         10.00         3.61         3.70         3.44           589479         AA167129         4.22         15.14         3.58         3.96         3.72           256260         H94617         1.40         4.95         3.54         3.05         3.96           256260         H94617         1.40         4.95         3.54         3.05         3.73           627273         AA191461         1.01         3.54         3.52         3.59         3.73           160436         AA488445         5.65         19.64         3.47         3.47         3.29           73139         AA448394         0.19         0.64         3.42         3.04         4.21           25553         R93309         0.72         2.45         3.41         3.01         3.43           412949         AA485773         1.36	-	0200	N57526	0.69	2.51	3.64	4.05	3.12	9 (	
136744         R.35230         2.00         7.22         3.52         3.26           487425         AA046523         2.77         10.00         3.61         3.70         3.44           589479         AA157129         4.22         15.14         3.58         3.96         3.72           256260         H94617         1.40         4.95         3.54         3.05         3.96           256260         H94617         1.40         4.95         3.54         3.05         3.96           25626         H94617         1.01         3.54         3.52         3.59         3.73           627273         AA191481         1.01         3.54         3.52         3.59         3.73           1604596         AA3962230         0.80         2.80         3.49         3.29         3.84           781339         AA448394         0.19         0.64         3.42         3.04         4.21           275553         R93309         0.72         2.45         3.41         3.01         3.43           412949         AA485773         5.81         19.81         3.41         3.01         3.43           412944         AA421273         1.36         4.57		27.50	A A 453870	3.81	14.21	3.64	3.48	4.25	3.19	•
176.744         K-35/30         2.00         3.61         3.70         3.44           487425         AA046523         2.77         10.00         3.61         3.70         3.44           589479         AA157129         4.22         15.14         3.58         3.96         3.72           256250         H94617         1.40         4.95         3.54         3.05         3.96           256250         H94617         1.40         4.95         3.54         3.05         3.96           256250         H94617         1.01         3.54         3.52         3.59         3.73           1604596         AA396230         0.80         2.80         3.49         3.29         3.84           843224         AA48845         5.65         19.64         3.47         3.47         3.36           27553         R93309         0.72         2.45         3.41         3.01         3.43           840394         AA485773         5.81         19.81         3.41         3.01         3.43           412849         AA421273         1.36         4.57         3.37         3.82         3.16           31044         AA421273         1.36         4.57		9	0.0000	6 6	4 3 3	3,52	3.22	3,26	4.37	က
487425         AA046523         2.77         10.00           589479         AA157129         4.22         15.14         3.58         3.96         3.72           256260         H94617         1.40         4.95         3.54         3.55         3.96         3.96           256260         H94617         1.40         4.95         3.54         3.52         3.59         3.73           627273         AA191481         1.01         3.54         3.52         3.59         3.73           1604596         AA48645         5.65         19.64         3.47         3.47         3.36           731339         AA448394         0.19         0.64         3.42         3.04         4.21           275553         R93309         0.72         2.45         3.41         3.01         3.43           840384         AA485773         5.81         19.81         3.41         3.01         3.43           412849         AA421273         1.36         4.57         3.37         3.82         3.16           31044         AA421273         1.36         4.57         3.37         3.82         3.16		5/44	K35230	6	00 01	3.61	3.70	3.44	3.70	ĸ
589479         AA157129         4.22         19.14         5.05         3.96           256260         H94617         1.40         4.95         3.54         3.05         3.96           256260         H94617         1.01         3.54         3.52         3.59         3.73           1604596         AA996230         0.80         2.80         3.49         3.29         3.84           843224         AA488445         5.65         19.64         3.47         3.47         3.36           781339         AA448394         0.19         0.64         3.42         3.04         4.21           275653         R93309         0.72         2.45         3.41         3.01         3.43           840384         AA485773         5.81         19.81         3.41         3.01         3.43           412849         AAA21273         1.36         4.57         3.37         3.82         3.16           731044         AA421273         1.36         4.57         3.37         3.82         3.16		37425		11.7	9.5	. e	3.96	3.72	3.07	က
255260         H94617         1.40         4.95         3.54         3.03         3.03           627273         AA191481         1.01         3.54         3.52         3.59         3.73           1604596         AA4986230         0.80         2.80         3.49         3.29         3.84           843224         AA488445         5.65         19.64         3.47         3.47         3.36           781339         AA448394         0.19         0.64         3.42         3.04         4.21           275653         R93309         0.72         2.45         3.41         3.42         3.22           440384         AA485773         5.81         19.81         3.41         3.01         3.43           412849         AAA707840         1.07         3.62         3.39         3.46         3.12           731044         AA421273         1.36         4.57         3.37         3.82         3.16		<b>39479</b>		77	4.0.	3	90 6	40 6	3.62	က
627273         AA191461         1.01         3.54         3.52         3.59         3.19           1604586         AA986230         0.80         2.80         3.49         3.29         3.84           843224         AA488445         5.65         19.64         3.47         3.47         3.36           781339         AA448394         0.19         0.64         3.42         3.04         4.21           275553         R93309         0.72         2.45         3.41         3.42         3.22           840384         AA485773         5.81         19.81         3.41         3.01         3.43           412949         AA707840         1.07         3.62         3.39         3.46         3.12           731044         AA421273         1.36         4.57         3.37         3.82         3.16	440 25	56260		<del>6</del> .	4.95	40.0	000	} <b>:</b>	3.24	n
1604596         AA996230         0.80         2.80         3.49         3.29         3.84           843224         AA488445         5.65         19.64         3.47         3.47         3.36           781339         AA448394         0.19         0.64         3.42         3.04         4.21           275653         R93309         0.72         2.45         3.41         3.42         3.22           840384         AA485773         5.81         19.81         3.41         3.01         3.43           412849         AA707840         1.07         3.62         3.39         3.46         3.12           731044         AA421273         1.36         4.57         3.37         3.82         3.16		27273		1.01	3.54	3.52	3.59	2,5	, « ; «	•
843224 AA488445 5.65 19.64 3.47 3.47 3.36 781339 AA448394 0.19 0.64 3.42 3.04 4.21 275653 R93309 0.72 2.45 3.41 3.42 3.22 840384 AA485773 5.81 19.81 3.41 3.01 3.43 412949 AA707840 1.07 3.62 3.39 3.46 3.12 731044 AA421273 1.36 4.57 3.37 3.82 3.16	•	04596		0.80	2.80	3.49	3.29	3.84	3.32	<b>~</b> (
245224     A4448394     0.19     0.64     3.42     3.04     4.21       275553     R93309     0.72     2.45     3.41     3.42     3.22       240384     AA485773     5.81     19.81     3.41     3.01     3.43       412949     AA707840     1.07     3.62     3.39     3.46     3.12       731044     AA421273     1.36     4.57     3.37     3.82     3.16		1327		5.65	19.64	3.47	3.47	3,36	3.59	· ·
275653 R93309 0.72 2.45 3.41 3.42 3.22 275653 R93309 0.72 2.45 3.41 3.01 3.43 840384 AA485773 5.81 19.81 3.41 3.01 3.43 412949 AA707840 1.07 3.62 3.39 3.46 3.12 731044 AA421273 1.36 4.57 3.37 3.82 3.16	•		•	pt 0	990	3.42	3.04	4.21	3.02	<b>6</b> 3
273553 K93309 0.12 2.75 840384 AA485773 5.81 19.81 3.41 3.01 3.43 412949 AA707840 1.07 3.62 3.39 3.46 3.12 731044 AA421273 1.36 4.57 3.37 3.82 3.16		2	•	, c	2.45	3.41	3.42	3.22	3.60	m
840384 AA485773 5.81 19.51 5.31 412949 AA707840 1.07 3.62 3.39 3.46 3.12 731044 AA421273 1.36 4.57 3.37 3.82 3.16		75653	H33309	<b>4</b> 5	7. 4	141	3.01	3.43	3.80	က
412949 AA707840 1.07 3.02 3.33 3.82 3.16	_	<del>6</del> 384		, i	10.61		3.46	3.12	3.59	က
731044 AA421273 1.36 4.57 3.3/ 3.02	•	12949	•	1.07	3.62	5.03	) r	3.5	3.13	ຕ
		31044	AA421273	1.36	4.57	3.37	3.82	2	<u>:</u> ;	

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Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

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<del>2</del> 4	501731	•	2.15	6.95	3.23	3.08	3.55	3.05	က
450	1640931	A1024401	0.21	0.66	3.13	3.14	3.07	3.17	က
451	299559		9.0	9.83	991.80	1.38	2966.06	7.96	~
452	365551	`	0.0	8.39	838.64	1.96	2506.90	7.05	7
453	42627	R60995	0.01	9.14	660.46	2:35	61.14	1917.92	~
454	277327		0.01	9.00	613.07	1.58	1831.96	5.67	7
455	773540		0.01	5.08	508.28	2.41	1516.35	6.10	7
456	12/1863		9.0	18.56	445.85	3.24	1331.89	2.42	7
457	754291		.00	¥.	433.98	2.48	1294.73	4.72	8
458	290213		0.01	3.94	393.88	0.73	1171.33	9.57	7
459	781157		0.0	2.83	283.26	1.00	841.07	7.73	7
460	781036		0.01	1.70	228.86	10.58	674.66	1.35	7
461	784216	AA446866	0.0	1.58	157.55	1.61	467.71	3.31	7
462	1377071		0.01	<del>7.</del> 2.	154.10	3.95	457.34	1.00	7
463	609935		0.02	2.74	114.55	1.41	338.20	4.03	7
48	203878		0.02	1.93	95.78	0.80	140.40	146.14	7
465	815794		0.08	6.83	85.65	0.89	150.46	105.59	2
466	347182		0.01	0.82	81.58	1.42	229.57	13.76	7
467	611373		0.01	0.71	70.86	2.42	205.80	4.35	~
468	746152		90.0	5.16	65.49	0.87	95.72	99.88	7
469	395708		0.01	0.83	60.41	6.88	173.62	0.73	7
470	684879		0.05	2.87	58.88	1.03	51.92	123.70	71
471	344589		0.22	12.41	56.02	78.84	88.50	0.72	7
472	562811		. 0.01	0.28	53.90	5.36	154.39	1.93	8
473	30207		9.0	2.21	52.11	0.54	150.40	5.39	7
474	1358266		9.08	3.95	51.39	1.05	149.73	3.40	7
475	757435		1.20	60.94	50.82	144.61	1.37	6.47	7
476	757143		0.19	9.88	50.76	4.68	145.07	2.54	7
477	33827		0.02	1.10	50.72	0.63	139.36	12.17	7
478	825740		0.26	13.02	49.67	0.91	56.58	91.51	8
479	813748		0.01	0.49	49.10	3.38	142.92	1.00	7
480	1505360		0.26	12.64	47.88	137.81	2.72	3.10	7
481	785849		0.01	0.60	45.99	2.24	128.30	7.45	~
482	43101	R59936	0.0	0.46	45.56	1.92	130.70	4.06	7
483	714498	AA292429	0.02	0.87	44.45	4.84	125.65	2.84	7
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Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

743362 1435339 28422 530608 774446 428828 196544 592928 773083 37823 37823 49858 743189 167205 53039 771294 167205 530900 280699 283379 283379 283379 283315 357884 172517 251806 80643 825606	3	811121	AA485688	0.01	0.44	44.03	3.52	127.57	00.1	۷ (
14.35239         AA857748         0.01         0.38         37.79         5.72         106.49           2.8422         R 14230         0.46         17.33         37.42         0.83         107.04           2.5066         AA071089         0.01         1.59         38.23         1.89         107.04           7.2466         AA40120         0.01         0.38         38.23         1.07         100.05           7.2862         AA40120         0.01         0.35         35.29         1.77         100.05           5.9928         AA40507         0.22         8.56         34.73         0.93         52.03           7.7308         AA42507         2.25         73.55         32.64         1.12         96.53           7.7308         AA42507         2.25         73.55         32.64         1.12         96.53           7.7308         AA42507         0.03         0.03         0.03         0.01         0.03         0.03           7.7308         AA42507         0.03         0.07         3.04         1.02         0.01           7.7308         AA43502         0.03         0.12         0.26         3.26         1.77         10.02	. 10	743362	AA400385	9.04	1.89	43.31	2.77	123.35	3,82	C1 1
28-62.2         R14230         0.46         17.33         37.42         0.83         107.04           530606         AAdrilog         0.01         0.38         35.23         3.89         107.04           77444         AAdrilog         0.01         0.35         35.29         1.20         103.0           77446         AAdrilog         0.01         0.35         35.29         1.77         100.36           592924         ARISON         0.03         0.35         35.29         1.77         100.36           592924         ARISON         0.18         6.00         33.64         1.12         96.53           77303         AAASSO         0.25         73.55         32.64         1.02         96.53           37823         AAASSO         0.03         1.12         32.46         1.79         96.53           37824         AAASSO         0.03         0.10         0.30         0.17         96.33           48956         AAAGSOS         0.02         0.07         0.30         0.07         1.00         7.97           48956         AAGSOS         0.01         0.30         0.30         0.77         96.38           16720         AAGSOS		1435339	AA857748	0.01	0.38	37.79	5.72	105.49	2.18	7
530608         AAAF01089         0.01         0.38         96.23         3.89         103.70           774446         AA446120         0.05         1.69         36.22         1.20         103.43           428828         AAA05224         0.01         0.35         36.29         1.77         100.85           196544         R91666         0.25         8.66         34.73         0.93         52.03           573232         AAA15377         0.18         6.00         3.26         1.17         96.63           773083         AAA62307         0.22         73.55         3.246         1.79         96.83           773083         AAA60434         0.02         0.70         30.56         2.89         0.12           743189         AAA6052         0.22         7.35         3.246         1.79         96.33           81586         AAA6052         0.02         0.70         30.56         2.89         0.12           81586         AAA6052         0.02         0.70         30.56         1.77         34.54           81586         AAA6052         0.02         0.70         30.66         2.89         0.12           81586         AAA6052		28422	R14230	0.46	17.33	37.42	0.83	107.04	4.40	2
774446         AAA46120         0.05         1.69         36.22         1.20         103.43           428828         AA005254         0.01         0.35         35.29         1.77         100.055           196544         R91566         0.25         8.56         34.73         0.93         52.03           592328         AA159179         0.18         6.00         32.64         1.12         96.63           77303         AA425307         2.25         73.55         32.46         1.12         96.63           77303         AA425307         2.25         73.56         1.79         96.53           74188         AA42602         0.03         2.02         32.46         1.12         96.63           81586         AA6903         0.03         1.12         32.46         1.12         96.53           81586         AA6805         0.03         0.07         0.00<		530608	AA071089	0.01	0.38	36.23	3.89	103.70	<del>.</del> :	61 (
228828         AAODGESA4         0.01         0.35         35.29         1.77         100.85           196544         R91566         0.25         8.56         34.73         0.93         52.03           59238         AA159179         0.18         6.00         33.64         1.12         96.53           773083         AA425307         2.25         73.55         32.64         1.12         96.53           773083         AA425307         2.25         73.55         32.64         1.12         96.38           773083         AA401434         0.03         2.62         31.64         1.12         96.38           81858         AA401434         0.03         0.70         30.56         2.89         61.70           48056         AA401434         0.02         0.70         30.66         1.00         7.97           48056         AA401434         0.02         0.70         30.06         1.00         7.97           48056         AA401434         0.01         0.20         2.76         2.83         6.17         7.25           48056         AA401434         0.01         0.20         30.06         1.77         3.45           48057		774446	AA446120	0.05	1.69	36.22	1.20	103.43	<del>4</del> 2	7
198544         R91566         0.25         8.56         34.73         0.83         52.03           592328         AA159179         0.18         6.00         33.64         1.12         96.53           773083         AA425307         2.25         73.55         32.64         1.02         96.53           773083         AA425307         2.25         73.55         32.64         1.02         96.53           49858         H29290         0.08         2.62         31.22         89.38         91.20           49858         H29290         0.08         2.62         31.22         89.38         91.20           48958         AA40044         0.02         0.70         30.56         2.89         91.20           48958         AA400502         0.21         5.88         27.65         1.74         98.44           48958         AA40302         0.01         0.30         27.63         1.74         3.45           48958         AA43524         0.01         0.20         27.63         1.74         70.25           48958         AA43524         0.01         0.20         27.63         1.74         70.25           71026         AA43524 <t< td=""><td></td><td>428828</td><td>AA005254</td><td>0.01</td><td>0.35</td><td>35.29</td><td>1.71</td><td>100.95</td><td>3.14</td><td>2</td></t<>		428828	AA005254	0.01	0.35	35.29	1.71	100.95	3.14	2
929232         AA459179         0.18         6.00         33.64         1.12         96.63           773083         AA42507         2.25         73.55         32.64         1.12         95.25           773083         AA42507         2.25         73.55         32.64         1.79         99.38           773083         AA42507         2.25         73.55         32.46         1.79         99.38           743169         AA40134         0.02         0.70         30.56         2.89         0.12           743169         AA405022         0.22         0.70         30.56         0.77         34.54           815861         AA405022         0.22         0.70         30.35         0.77         34.54           439565         AA40322         0.01         0.30         30.08         1.00         7.97           43957         AA403274         0.01         0.20         27.63         1.51         76.38           77124         AA425158         0.01         0.27         27.13         0.74         70.25           776500         AA425158         0.01         0.20         27.04         8.33         1.35           77650         AA425158		196544	R91566	0.25	8.56	34.73	0.93	52.03	51.22	7
733083         AA425307         2.25         73.55         32.64         1.02         93.52           37823         RA425307         2.25         1.12         32.46         1.79         99.38           49B65         H29290         0.08         2.62         31.22         87.92         0.12           743169         H29230         0.08         2.62         31.22         87.92         0.12           815861         AA48602         0.28         0.29         30.36         0.77         34.54           89585         AA409359         0.01         0.30         30.08         1.00         7.97           48956         AA43936         0.21         5.88         27.65         1.51         34.54           5003         R15740         0.01         0.20         27.63         1.51         76.38           71124         AA443624         0.01         0.27         27.13         1.21         76.38           71124         AA445624         0.01         0.27         27.13         1.21         7.20           71841         H50655         0.53         14.38         26.99         2.23         1.35           71891         H50655         0.53<		59292R	AA159179	0.18	6.00	33.64	1.12	96.63	3.18	7
37823         R59473         0.03         1.12         32.46         1.79         89.38           49858         H29290         0.08         2.62         31.22         87.92         0.12           743189         AA401434         0.02         0.70         30.56         2.89         81.20           815861         AA408022         0.28         7.89         30.35         0.77         34.54           815865         AA408022         0.26         7.89         30.35         0.77         34.54           489656         AA40332         0.21         5.88         27.65         1.51         76.38           55039         R45740         0.01         0.20         27.73         3.44         77.27         3.44           771294         AA4450274         0.01         0.27         27.13         1.27         70.25           771294         AA4450274         0.01         0.20         27.73         27.37         3.84           771294         AA4450274         0.01         0.27         27.13         1.27         2.05           78000         AA443024         0.01         0.27         27.13         1.27         2.05           78001         <		773083	AA425307	2.25	73.55	32.64	1.02	93.52	3.37	71
49B56         H25200         0.08         2.62         31.22         87.92         0.12           743189         A4401434         0.02         0.70         30.56         2.89         81.20           743189         A4401434         0.02         0.70         30.56         2.89         81.20           815861         A4485052         0.26         7.89         30.35         0.77         34.54           489656         A4003329         0.01         0.30         30.08         1.00         7.97           489656         R40339         0.21         5.88         27.63         1.51         76.38           489656         R40339         0.01         0.27         27.34         0.74         70.25           771234         A443624         0.01         0.27         27.34         0.74         70.25           771234         A443624         0.01         0.27         27.13         1.21         6.904           771234         A443624         0.01         0.20         2.73         27.34         0.74         70.25           280699         N47445         0.65         0.59         0.69         2.78         6.904           28069         N		47.873	P50473	0.03	1,12	32.46	1.79	89.38	6.23	7
435183         AA401434         0.02         0.70         30.56         2.89         81.20           4351861         AA485052         0.26         7.89         30.35         0.77         34.54           485656         AA485052         0.26         7.89         30.35         0.77         34.54           485656         AA403329         0.01         0.30         30.08         1.00         7.97           485656         AA60324         0.28         16.16         27.63         1.51         76.38           53029         R15740         0.01         0.27         27.34         0.74         70.25           771234         AA443524         0.01         0.27         27.34         0.74         70.25           771234         AA443524         0.01         0.27         27.34         0.74         70.25           184318         H50655         0.53         14.38         26.99         0.69         27.83           184318         H50655         0.53         14.15         25.90         2.28         66.04           280699         M7445         0.65         14.15         25.90         2.28         67.88           28069         M4745	<b>.</b> 11	4985B	H29290	0.08	2.62	31.22	87.92	0.12	5.61	2
445652         0.26         7.89         30.35         0.77         34.54           480565         AAA95052         0.26         7.89         30.35         0.77         34.54           480565         AAA93539         0.01         0.30         30.08         1.00         7.97           480563         AA49359         0.21         5.88         27.65         1.51         76.38           59039         Regass         0.23         27.34         0.74         70.25           77124         AA443624         0.01         0.27         27.33         1.35         9.84           77124         AA443624         0.01         0.27         27.34         0.74         70.25           77124         AA443624         0.01         0.20         27.03         26.99         0.69         27.83           77124         AA443624         0.01         0.20         27.04         8.33         1.35           77124         AA443624         0.01         0.20         27.04         8.33         1.35           78009         AA435188         0.01         0.20         27.04         8.33         1.75           2806         AA0018         0.01         0.25		43000	087674		0.70	30.56	2.89	81.20	2.60	7
489655         AA099359         0.01         0.03         0.06         7.97           489655         AA099359         0.01         0.30         0.06         7.97           167205         R49439         0.01         0.27         27.63         72.37         9.84           53039         R15740         0.01         0.27         27.34         0.74         70.25           795503         R445624         0.01         0.27         27.34         0.74         70.25           77124         AA440524         0.01         0.27         27.34         0.74         70.25           77124         AA440524         0.01         0.20         27.04         8.33         1.35           77124         AA440524         0.01         0.20         27.04         8.33         1.35           77129         AA440524         0.01         0.20         27.04         8.33         1.35           77129         AA425158         0.01         0.20         27.04         8.33         1.35           280699         NA7445         0.65         14.15         25.06         66.04           27605         RA00498         0.01         0.25         25.21         2.2	n :	801647	AA401434	20.0 0.04	7.89	30.35	0.77	25.52	55.73	7
4030309         COLOR         <		100010	A A COCCEO	2	030	30.08	1.00	7.97	81.28	2
167205         R80934         0.58         16.16         27.63         72.37         9.84           53039         R15740         0.01         0.27         27.34         0.74         70.25           771294         AA402274         0.01         0.27         27.13         1.21         69.04           771294         AA403524         0.01         0.20         27.04         8.33         1.36           164318         H50655         0.63         14.38         26.99         0.69         27.83           769000         AA425158         0.01         0.26         26.26         5.77         72.00           280699         N47445         0.65         14.15         25.90         2.28         11.75           280799         NA7445         0.65         14.15         25.90         2.28         11.75           280795         NA7445         0.65         14.15         25.90         2.28         66.04           280795         NA7445         0.02         0.39         25.57         2.05         68.00           280795         AA401496         0.07         1.90         25.51         2.29         67.88           2316         AV94486		76707	60033003	12.0	88	27.65	1.51	76.38	5.06	7
53039         R15740         0.01         0.27         27.34         0.74         70.25           796549         AA460274         0.01         0.27         27.13         1.21         69.04           771294         AA460274         0.01         0.20         27.04         8.33         1.36           194318         H50655         0.53         14.38         26.99         0.69         27.83           789000         AA425158         0.01         0.26         26.26         5.77         72.00           280699         NA7445         0.65         14.15         26.99         0.69         27.83           280699         NA7445         0.65         14.15         26.90         2.28         11.75           280699         NA7445         0.65         14.15         26.90         2.28         11.75           280699         NA7445         0.02         0.39         25.65         0.59         66.04           27605         R40018         0.07         1.90         25.57         2.05         66.04           27805         AA400486         0.01         0.25         25.1         2.29         67.88           251806         H36647 <td< td=""><td>» c</td><td>467205</td><td>D00004</td><td>0.58</td><td>16.16</td><td>27.63</td><td>72.37</td><td>9.84</td><td>0.70</td><td>7</td></td<>	» c	467205	D00004	0.58	16.16	27.63	72.37	9.84	0.70	7
796549         AAA45024         0.01         0.27         27.13         1.21         69.04           771294         AAA43024         0.01         0.20         27.04         8.33         1.36           1961318         H50555         0.53         14.38         26.99         0.69         27.83           769000         AA425158         0.01         0.26         26.26         5.77         72.00           280699         N47445         0.65         14.15         25.90         2.28         11.75           280699         N47445         0.65         14.15         25.90         2.28         17.50           280699         N47445         0.02         0.39         25.57         2.06         68.60           27605         R40018         0.07         1.30         25.57         2.05         68.60           27605         R40018         0.07         0.25         25.17         2.29         66.04           27605         R40018         0.07         0.25         25.17         2.05         66.04           27816         AA701996         0.07         0.25         25.17         2.86         67.40           25316         H19624		6303	D45740	100	0.27	27.34	9.74	70.25	11.03	64
71724         AA443624         0.01         0.20         27.04         8.33         1.36           194318         H50655         0.53         14.38         26.99         0.69         27.83           769000         AA425158         0.01         0.26         26.26         5.77         72.00           280699         N47445         0.65         14.15         25.90         2.28         11.75           280599         N47445         0.05         0.39         25.65         0.59         66.04           283379         N52767         0.02         0.39         25.65         0.59         66.04           27605         AA070437         0.01         0.25         25.71         2.05         68.60           436121         AA701996         0.02         0.60         25.17         3.88         71.30           436121         AA701996         0.01         0.25         25.17         3.86         71.73           435784         W94486         0.01         0.25         24.95         2.37         41.17           17517         H19626         1.00         24.91         24.95         2.27         26.84           251805         AA490945		705549	A A S C C C A A A A A A A A A A A A A A	0.01	0.27	27.13	1.21	P0.69	11,15	7
194318         H50655         0.53         14.38         26.99         0.69         27.83           769000         AA425158         0.01         0.26         26.26         5.77         72.00           280699         N47445         0.65         14.15         25.90         2.28         11.75           280379         N52767         0.02         0.39         25.65         0.59         66.04           27605         R40018         0.07         1.90         25.57         2.05         68.60           530958         AA70437         0.01         0.25         25.21         2.29         68.60           530958         AA70496         0.02         0.60         25.17         3.88         71.30           436121         AA70196         0.02         0.60         25.17         3.88         71.30           53316         H19826         0.01         0.25         25.17         3.88         71.30           53318         W94486         0.01         0.25         24.95         2.32         41.17           1751         H19826         1.00         25.495         2.486         2.77         26.82           25480         H9647         0.01		771794	A A 443624	100	0.20	27.04	8.33	1.35	71.43	2
789000         AA425158         0.01         0.26         26.26         5.77         72.00           280699         N47445         0.65         14.15         25.90         2.28         11.75           280699         N47445         0.65         14.15         25.90         2.28         11.75           281379         N62767         0.02         0.39         25.65         0.59         66.04           27605         R40018         0.07         1.90         25.57         2.05         68.60           436121         AA701996         0.07         0.25         25.11         2.29         67.88           436121         AA701996         0.02         0.60         25.17         3.88         71.30           435181         0.01         0.25         25.10         2.87         64.40           357884         W94486         0.01         0.25         24.95         2.37         41.17           172517         H19626         1.00         24.91         24.95         2.27         26.84           251806         H96647         0.01         0.24         24.50         1.76         60.46           824025         AA490045         0.31 <t< td=""><td></td><td>194318</td><td>H50655</td><td>0.53</td><td>14.38</td><td>26.99</td><td>0.69</td><td>27.83</td><td>52.45</td><td>7</td></t<>		194318	H50655	0.53	14.38	26.99	0.69	27.83	52.45	7
280699         NA7445         0.65         14.15         25.90         2.28         11.75           280699         NA7445         0.65         14.15         25.65         0.59         66.04           280699         NA7445         0.02         0.39         25.65         0.59         66.04           27605         R40018         0.07         1.90         25.57         2.05         68.60           436121         AA701996         0.07         0.60         25.17         3.86         71.30           436121         AA701996         0.02         0.60         25.17         3.86         71.30           357884         W94486         0.01         0.25         24.95         2.37         41.17           172517         H19626         1.00         2.49         2.37         41.17           251806         H96647         0.01         0.24         24.50         1.76         60.48           80643         157803         1.09         26.54         24.25         0.44         9.54           824025         AA490045         0.31         7.41         24.24         0.95         28.65           838862         AA490044         0.28 <td< td=""><td>,</td><td>769000</td><td>AA425158</td><td>0.01</td><td>0.26</td><td>26.26</td><td>5.77</td><td>72.00</td><td>9.</td><td>7</td></td<>	,	769000	AA425158	0.01	0.26	26.26	5.77	72.00	9.	7
283379         N52767         0.02         0.39         25.65         0.59         66.04           27605         R40018         0.07         1.90         25.57         2.05         68.60           530956         AA070437         0.01         0.25         25.21         2.29         67.88           436121         AA701996         0.02         0.60         25.17         3.86         71.30           53315         R15813         0.01         0.25         24.95         2.87         64.40           35784         W94486         0.01         0.25         24.95         2.32         41.17           172517         H19826         1.00         24.91         24.86         2.27         26.82           251806         H96647         0.01         0.24         24.50         1.76         60.48           80643         T57803         1.09         26.54         24.25         0.44         9.54           824025         AA490045         0.31         7.41         24.24         0.95         28.65           82866         AA490044         0.28         6.50         23.31         0.87         64.54           45         AA490044         0.28		280699	NAZAAS	0.55	14,15	25.90	2.28	11.75	63.66	2
27605         RA40018         0.07         1.90         25.57         2.05         68.60           530956         AA070437         0.01         0.25         25.21         2.29         67.88           436121         AA701996         0.02         0.60         25.17         3.88         71.30           53315         R15813         0.01         0.25         25.10         2.87         64.40           35784         W94486         0.01         0.25         24.95         2.32         41.17           172517         H19626         1.00         24.91         24.86         2.27         26.82           251606         H96647         0.01         0.24         24.50         1.76         60.48           80643         T57803         1.09         26.54         24.25         0.44         9.54           824025         AA490945         0.31         7.41         24.24         0.95         28.65           825606         AA490044         0.28         6.50         23.31         0.87         64.54           838882         AA490044         0.28         6.50         23.31         0.87         64.54           15         15         15 <td></td> <td>281379</td> <td>N52767</td> <td>0.02</td> <td>0.39</td> <td>25.65</td> <td>0.59</td> <td><b>66.04</b></td> <td>10.31</td> <td>8</td>		281379	N52767	0.02	0.39	25.65	0.59	<b>66.04</b>	10.31	8
530958         AA070437         0.01         0.25         25.21         2.29         67.88           436121         AA701996         0.02         0.60         25.17         3.88         71.30           53315         R15813         0.01         0.25         25.10         2.87         64.40           35784         W94486         0.01         0.25         24.95         2.32         41.17           172517         H19626         1.00         24.91         24.86         2.27         26.82           251806         H96647         0.01         0.24         24.50         1.76         60.48           80643         757803         1.09         26.54         24.25         0.44         9.54           825606         AA504625         0.31         7.41         24.24         0.95         28.65           838862         AA490044         0.28         6.50         23.31         0.87         64.54           15         15         15         16         16         16         16         16         16		27505	RADOTR	0.07	1.90	25.57	2.05	68.60	6.05	2
436121         AA701996         0.02         0.60         25.17         3.88         71.30           53315         R15813         0.01         0.25         25.10         2.87         64.40           357884         W94486         0.01         0.25         24.95         2.32         41.17           175517         H19626         1.00         24.91         24.86         2.27         26.82           251806         H9647         0.01         0.24         24.50         1.76         60.46           80643         757803         1.09         26.54         24.25         0.44         9.54           824025         AA490945         0.31         7.41         24.24         0.95         28.65           825606         AA504625         0.15         3.51         23.68         0.87         21.91           839882         AA490044         0.28         6.50         23.31         0.87         64.54           15         15         16         176         66.54         176         64.54		530958	AA070437	0.0	0.25	25.21	2.29	67.88	5.46	7
53315         R15813         0.01         0.25         25.10         2.87         64.40           35784         W94486         0.01         0.25         24.95         2.32         41.17           172517         H19626         1.00         24.91         24.86         2.27         25.82           251806         H9647         0.01         0.24         24.50         1.76         60.46           80543         T57803         1.09         26.54         24.25         0.44         9.54           824025         AA490945         0.31         7.41         24.24         0.95         28.65           825606         AA504625         0.15         3.51         23.68         0.87         21.91           839882         AA490044         0.28         6.50         23.31         0.87         64.54           15         15         15         16         16         16         16         16		436434	AAYN4GGE	200	0.60	25.17	3.88	71.30	9.34	8
357884         W94486         0.01         0.25         24.95         2.32         41.17           172517         H19826         1.00         24.91         24.86         2.27         26.82           251806         H9647         0.01         0.24         24.50         1.76         60.48           80643         T57803         1.09         26.54         24.25         0.44         9.54           824025         AA490945         0.31         7.41         24.24         0.95         28.65           825606         AA504625         0.15         3.51         23.68         0.87         21.91           839882         AA490044         0.28         6.50         23.31         0.87         64.54           15         15         16         16         16         16         16		53315	R15813	100	0.25	25.10	2.87	64.40	8.05	7
172517         H19826         1.00         24.91         24.86         2.27         26.82           251806         H96647         0.01         0.24         24.50         1.76         60.48           80643         T57803         1.09         26.54         24.25         0.44         9.54           824025         AA490945         0.31         7.41         24.24         0.95         28.65           825606         AA504625         0.15         3.51         23.68         0.87         21.91           839882         AA490044         0.28         6.50         23.31         0.87         64.54           15         15         15         15         15         15	- ^	357884	W94486	0.0	0.25	24.95	2.32	41.17	31.35	7
251806 H96647 0.01 0.24 24.50 1.76 60.48 80643 T57803 1.09 26.54 24.25 0.44 9.54 8.54 82506 AA504625 0.31 7.41 24.24 0.95 28.65 82506 AA504625 0.15 3.51 23.68 0.87 21.91 839882 AA490044 0.28 6.50 23.31 0.87 64.54 15		172517	H19826	00.	24.91	24.86	2.27	26.82	45.50	7
80643 T57803 1.09 26.54 24.25 0,44 9.54 824025 AA490945 0.31 7.41 24.24 0.95 28.65 825606 AA504625 0.15 3.51 23.68 0.87 21.91 839882 AA490044 0.28 6.50 23.31 0.87 64.54	. ~	251806	H96647	0.0	0.24	24.50	1.76	60.48	11.25	2
824025 AA490945 0.31 7.41 24.24 0.95 28.65 825606 AA504625 0.15 3.51 23.68 0.87 21.91 839882 AA490044 0.28 6.50 23.31 0.87 64.54	- w	R0543	157803	1.09	26.54	24.25	4.	9.54	62.76	7
825608 AA504625 Q.15 3.51 23.68 0.87 21.91 839882 AA490044 0.28 6.50 23.31 0.87 64.54 15		824025	AA490945	0.31	7.41	24.24	0.95	28.65	43.13	2
839862 AA490044 0.28 6.50 23.31 0.87 64.54 15		825606	AA504625	D 15	3.51	23.68	0.87	21.91	48.25	2
		839882	AA490044	0.28	6.50	23.31	0.87	64.54	4.51	7
			!			5				

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

8	8	7	64	8	7	7	~	7	7	~	<b>C4</b>	2	2	7	7	8	73	2	7	7	7	~	7	7	7	8	7	7	7	2	7	7	7	2	
25.34	0.73	53.51	54.20	23.77	3.87	3.43	17.48	3.78	28.27	20.17	50.17	45.41	25.26	49.74	46.16	1.08	5.54	6.22	5.54	16.22	8.77	127	12.21	34.18	7.22	5.65	28.36	25.15	3.27	0.57	15.13	18.34	3.12	6.87	
42.67	61.72	9.53	10.05	1.50	58.79	57.89	42.76	0.20	30.54	38.72	7.58	10.31	30.81	1.00	1.93	14.31	47.06	45.91	45.44	34.15	40.61	3.64	37.48	1.51	39.78	40.90	16.52	21.09	41.42	41.20	28.43	26.10	38.63	0.27	
1.00	3.92	2.34	0.88	39.83	0.72	0.93	0.93	56.76	1.29	1.00	1.10	2.79	2.35	7.38	9.69	41.41	2.02	1.21	1.87	1,97	2.37	48.76	1.87	14.65	2.86	2.84	2.63	0.69	1.59	3.92	1.53	0.42	2.32	36.64	
23.00	22.12	21.79	21.71	21.70	21.13	20.75	20.39	20.24	20.04	19.97	19.61	19.50	19.47	19.37	19.26	18.93	18.21	17.78	17.62	17.44	17.25	17.22	17.19	16.78	16.62	16.47	15.84	15.64	15.42	15.23	15.03	14.95	14.69	14.60	16
0.23	0.30	0.34	3.65	0.14	0.51	0.95	6.19	1.02	3.94	0.20	3.26	0.20	3.30	0.19	0.19	21.01	0.74	3.45	0.54	0.17	0.44	8.76	1.06	0.11	0.29	1.14	2.94	5.98	2.60	0.95	3.91	3.94	1.38	0.54	
0.01	0.0	0.02	0.17	0.01	0.02	0.05	0.30	0.05	0.20	0.01	0.17	0.01	0.17	9.0	0.01	1.1	9.	0.19	0.03	0.01	0.03	0.51	90.0	0.01	0.02	0.07	0.19	0.38	0.17	90.0	0.26	0.26	0.09	9.0	
170198	H88143	AA161097	AA491261	H10072	W95757	. N24848	AA255954	R43910	R95684	AA461508	AA520979	R86970	H28091	AA496022	AA456628	T65736	AA620747	AA487505	R60020	AA001376	AA489662	AA857809	AA857163	AA481481	N26559	N51297	AA504838	AA417713	AA206311	N73083	AA463445	H18932	R49592	AA035745	
81050	220655	591055	824270	46843	358316	267085	686552	32962	199337	795831	826256	197265	162310	759163	809583	80338	1049284	839048	42907	361899	823727	1475476	1410444	756629	266366	283070	825726	746232	647679	291947	811766	51599	37205	364885	
519	520	521	522	523	524	525	526	527	528	529	530	531	532	533	534	535	536	537	538	539	540	3	542	543	3	545	546	547	548	549	550	551	552	553	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	2	7	7	7		, ,	<b>7</b> (	7	7	7	7	7	2	. 2		• •	• (	7 4	2	7	7	7	2	7	7	2	7	7	2	7	2	7	8	2	
2.07	23.25	2.72	a S	4.61	3.01	5.10	27 68	3	12.18	18.27	29.19	26.50	7:37	25.70	28.39	2 23	4	2 4	97.4	0.58	<b>7</b> .	26.39	20.98	11.09	1.00	4.42	20.10	3.37	16.24	28.00	3.4	14.01	28.79	3.60	3.82	3.43	i
38.09	18.93	36.42 1	37,99	35.90	2.96	38.28	20.20	\$ :	26.44	21.58	8.82	1.54	9.78	12.28	8 17	20.04	7 77	31.49	31,18	14.55	29.00	0.39	14.19	23.87	29.43	2.25	14.95	30.06	17.13	4.48	30.95	20.54	0.92	28.84	27.04	28.86	) ) 
3.43	1.00	3.59	1.58	1.89	36.41	, a	0.68	2.99	2.42	0.64	2.08	10.97	30.80	0.65	88	3	<b>6</b> .,	1.33	2.47	22.57	1.18	10.40	1.96	2.11	6.29	, 29.85	1.00	2.42	2.08	2.89	0.81	0.58	4.71	1.92	2.83	96 -	3
14.53	14,40	14.24	14.20	14 13	7	5	14.09	13.89	13.68	13.50	13.36	13.00	12.97	12 88		10.21	12.79	12.64	12.63	12.56	12.41	12.39	12.38	12.36	12.24	12.17	12.02	11.95	11.82	11.78	11.74	11.71	11,47	11.45	11.23	2 7	77.11
5.51	140	0.43	7 7	, ,		12.46	2.19	4.17	1,31	3.95	8.41	80.0	35.0	) ; ;	2 1	8.32	0.71	1.99	0.33	0.13	4.99	44.0	0.12	3.72	0.12	18.67	1.69	0.52	23.53	0.12	0.12	8 0	21.0	0.40	? 9	04.7	2.42
ğ		5 6	3.5	7.0	60.0	0.88	0.16	030	0.10	0.29	<b>1 2 3</b>	0		3 3	5	0.65	9.06	0.16	9.0	0.01	0.40	0.78	2		50.0	53	0.14	40.0	8	5	5 6	5 6	3 6	2	3 8	8	22
36463044	***************************************	AA434388	AA453260	R40105	AA010224	N53380	AA699501	W90705	R37738	AA427749	LAEEAO	NEO-103	102133	AA035730	AA017379	AA911238	R43915	AA070435	H11051	AA453616	44437212	AAE20607	4400200	MU22490	750537	45154 AAARAR17	Pa2201	AA308324	44477227	18/61706	4446643	******	1101100	1987/00	G1081H	AA478952	AA609300
1155603	23000	0990//	795347	30082	430237	284001	436402	418004	26806	774004		284288	207407	30484	361122	1526789	33066	530954	47204	795446	757A7B	40,404	364353	70000	71674	1,071	105845	706707	730450	224404	106367	1989	101180	707005	20805	753984	1031561
į	g i	555	556	221	558	559	260	564	, Ç		3 3	<b>3</b> 4	8	28	567	568	569	570	7.5	: 8	, E	3 3		6/9	6		0 2		200	- ·	78C	2	8	582	586	287	588

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelia! Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

~	7	8	7	~	8	7	7	7	~	7	7	8	7	~	~	7	8	7	7	7	7	7	64	N	7	7	7	~	7	7	8	8	7	7	
16.34	7.83	27.27	27.95	13.80	<del>6</del> .8	4.84	16.62	10.20	.3.28	20.10	8.36	4.31	21.77	4.40	4.66	2.78	6.78	4:47	15.60	3.50	23.76	15.64	0.48	1.65	1.83	17.42	21.42	1.12	16.30	1.00	15.31	24.03	10.88	1.81	
16.85	22.78	4.45	1.00	17.96	26.77	25.57	15.37	20.21	27.39	10.22	1.00	25.25	8.99	25.88	1.37	23.39	21.63	24.49	14.48	24.58	6.31	14.23	7.21	17.15	25.02	11,85	9.09	20.18	12.15	24.20	13.69	0.98	17.62	19.71	
0.33	2.56	1,10	3.77	0.92	1.90	2.21	0.51	2.09	1.79	2.05	22.91	2.40	0.74	1.7	25.10	4.91	2.67	2.40	0.88	2.87	0.86	0.77	22.87	11.51	3,45	0.93	2.67	8.61	1.40	4.50	0.67	4.65	1.10	8.07	
11.17	11.06	10.94	10.91	10.90	10.89	10.87	10.84	10.83	10.82	10.79	10.76	10.65	10.50	10.47	10.38	10.36	10.36	10.35	10.32	10.32	10.31	10.21	10.19	10.11	10.10	10.07	10.05	9.97	9.95	9.80	9.89	9.89	9.87	9.86	18
0.34	0.95	1.32	0.11	3.90	0.36	1.12	6.12	0.31	3.13	9.32	0.11	7.96	8.09	7.51	18.95	12.67	4.0	0.21	0.87	4.59	4.21	5.55	9.47	0.12	8.21	2.53	16.05	2.42	2.29	0.10	4.94	45.45	3.82	0.09	
0.03	0.09	0.12	0.0	0.36	0.03	0.10	0.57	0.03	0.29	0.86	0.0	0.75	0.77	0.72	1.83	1,22	20.02	0.02	0.08	0.45	0.41	0.54	0.93	0.0	0.81	0.25	1.60	0.24	0.23	Q.01	0.50	4.60	0.39	0.01	
AA160670	AA427737	AA454222	N31808	H48070	AA453787	R66438	AA410434	N74131	N66900	N30751	AA680136	R39765	AA489073	AA463248	AA702803	AA421171	N93721	AA418852	AA281784	AA099148	H54451	N49389	AA412049	AA158035	AA775223	R94504	AA478036	AA626167	AA664007	AA430042	AA292226	R25377	N21633	R55220	
807728	770854	795526	271483	193333	813738	41869	753428	298417	287190	257955	433155	26164	824802	797055	448036	731095	307138	768043	712401	489509	203184	277621	729953	590640	868838	197651	740457	745248	855391	781110	725877	133136	268135	154720	•
OR A	280	591	592	593	594	585	298	597	598	685	009	601	602	603	908	605	909	607	808	609	610	611	612	613	614	615	616	617	618	619	920	621	622	623	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

																														•								
•		, (	79 (	7	2	7	2	7	7	. ~			4 (	J F	7 6	4 (	<b>v</b>	N C	7 (	7	7	7 (	7	7 (	7	Ν.	7	7	7	7	7	7	7	7	7	2		
47	2 .	6.03	3.25	5.79	18.58	4.67	6.28	3.39	85	08.0+	20.00	7	60.1		16.97	18.40	21.00	3.86	18.37	28.	15.08	14,44	21.78	23.74	9.79	17.71	5.45	13.51	14.35	21.12	14.67	5.38	9.46	1.10	4.17	16.60	·	
	23.34	22.72	24.64	22.17	9.16	23.22	21.91	24 53		8.7	30.61	3.74	3.03	11.85	7.00	40.7	5.32	23.58	1.12	21.39	1.54	12.53	1.86	1.00	16.72	7.70	19.73	13.04	0.36	1.00	10.87	20.24	15.22	13.45	21.38	) (2)	0:30	
!	5.59	2 2 2	1.50	1.35	1.31	1 13	 	9 6	0.0	1.67	2.05	24.42	23.58	1.15	10.45	8.70	<del>1</del> .82	0.66	8.59	4.86	11.43	0.97	4.26	3.14	1.35	2.30	2.37	0.81	12.50	5.08	1.65	<u>.</u> 2	2.44	12 54	5.4	24.1	10.05	
	9.80	9.80	9.79	77.6	89	0.00 FB 0	0.6	3.62	9.57	9.56	9.53	9.51	9.50	9.48	9.47	9.40	9.39	9.37	9.36	9.36	9.34	9.31	9.30	9.29	9.29	9.26	9.18	9.12	9.07	9.07	90.6	20.0	2		9.03	50 G	8.98	<u>0</u>
	0.21	6.27	2.73	0.21	F 0.7	9.6	0.10	6.12	4.60	0.16	0.77	6.70	12.98	1.40	0.09	0.09	203	3.84	0.08	0.32	90'0	0.22	0.09	0.09	6.35	4.10	0.87	244	0.26	60.0	12.5		3.02	20.01	11.71	4.35	17.81	
	0.02	0.64	0.28	200		0.62	0.0	0.64	0.48	0.02	0.08	0.70	1.37	0.15	0.01	0.01	0.53	0.41	0.0	500	3 6	0.02	100	0.0	890	77	0.10	7.2.0	500	6	5 6	2 6	9.4 9. i	1.1	1.30	0.48	1.98	
	AA464615	AA252080	D40478	071694	61/07/50	AA620528	AA447986	AA133590	AA487432	R89490	R46000	AAD18980	AA032221	AA219060	W16659	AA679180	AA13664	H11519	AA699926	MEASEA	A A 28 4 0 7 2	NKOG62	200000	AA045074	AAAEE343	A1047607	A101/00/	100000	RJCORY	0000000VVV	AA100090	NZ6515	AA192527	AA488175	AA172056	AA158346	AA620485	
	812994		20500	06000	100084	951305	782725	512116	841323		35804	362928	375682			866702	490959	4822B	738330	20001	24/040	284463	101107	487165	201704	187410	730882	700007	176861	200026	49055	52997	627541	877632	594758	591095	951241	
	824	, ,	3	979	/79	828	629	830	<u>8</u>	632	83	2	535	9 5	2 2	3 6	3 6	3 3	3 3	3	7 5	3 3	1	649	3	\$	8 8	2	8 3	ري و	652	653	654	655	999	657	658	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	7	7	7	7	7	8	7	7	8	2	7	64	N	8	7	7	8	ru	2	7	2	8	~	7	7	~	~	8	7	<b>~</b>	~	7	7	8	
5.40	11.94	8,45	14.43	8.30	18.10	0.61	3.16	7.50	3.36	14.35	18.86	17.99	19.50	11.27	12.36	17.21	2.28	16.24	19.27	2.96	13.09	11.26	0.50	18.07	3.68	9.48	4.38	17.47	5.36	13.46	18.54	8.40	8.68	3.15	•
20.84	2.96	17.16	2.08	17.44	6.64	21.88	21.71	16.58	21.85	1.00	0.52	6.50	3.76	12.46	11.61	6.16	18.57	6.27	0.41	13.18	1.67	11.14	19.58	0.56	20.61	13.54	17.84	2.94	18.34	B.34	4.66	14.68	13.94	18.08	
0.65	11.72	0.92	9.92	0.36	1.3 E.	3.59	1.16	1.87	0.60	10.33	6.22	0.78	1.96	1.45	1.30	1.81	4.15	2.41	5.22	8.74	96.6	2.23	4.39	5.83	0.11	1.22	1.99	3.70	0.38	2.25	0.81	0.92	<b>4</b> .06	2.43	
8.96	8.87	8.84	8.81	8.70	8.69	8.69	8.68	8.65	8.60	8.56	8.53	8.42	8.41	8.39	8.39	8.39	8.34	8.31	8.30	8.29	8.25	8.21	8.16	8.15	8.13	8.08	8.07	8.04	8.03	8.02	8.00	8.00	7.89	7.89	20
0.21	5.97	3.32	2.18	39.21	0.80	4.61	9.68	34.28	6.05	0.09	5.11	6.98	11.80	9.51	3.36	0.08	80.0	13.45	10.75	1.09	0.15	1,49	0.16	0.14	21.79	6.11	. 18.15	60:0	0.22	0.08	5.71	8.81	2.29	15.64	
0.02	29'0	0.38	0.25	4.51	60.0	0.53	1.12	3.98	0.59	0.01	0.60	0.83	1.40	1.13	0.40	0.01	0.0	1.62	1.30	0.13	0.02	0.18	0.02	0.02	2.68	0.76	2.00	0.01	0.03	0.01	0.71	1.10	0.29	1.98	
AA416970	W68220	H41203	H16772	AA629999	AA778646	W72671	AA609471	AA775863	R40373	AA496998	AA682321	N80764	T87515	W47364	AA504265	AA036974	AA132226	N74617	AA488324	R61372	N50806	AA058576	R47893	AA131239	AA447504	AA699427	AA701351	AA460961	AA457576	H94944	AA487674	H51765	AA284172	W90381	
730036	342640	176533	50227	884511	1048995	345761	743589	878544	28927	897546	462926	300615	115408	324717	825416	484535	566887	296429	842968	37980	280882	380884							838776		841620	193892	701625	417920	
629	660	661	662	663	664	665	999	299	668	699	670	671	229	673	674	675	576	<i>677</i>	678	679	680	189	682	683	684	685	989	687	688	689	069	69	692	693	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	~	2	ור	<b>J</b> (	8	~	~	7	7	8	7	2	8	8		, ,	• •	7 (	7 (	7	7	7	7	7	8	7	8	7	7	~		۰ ۲	• ^	• ^	•
2.87	3.76	6.01	18.64	000	3	07.61	4.90	9.28	1.00	17.46	9.03	29.	2.00	3.72	5.66	632	45.79		2.32	15.89	9.88	15.83	1.19	9.18	4.60	12.04	9.23	15.98	3.75	8.02	15.57	14 40	75.64	٠٠٠٠	C7'1	27.0L	12.41
16.63	1.50	15.21	3 O.	5 .	20.	7.43	17.32	12.63	14.11	3.87	41 89	14.72	9 20	15.57 C7. 24	40.05		12.70	5.74	17.38	<b>1</b> .8	12.36	5.39	18.13	0.92	15.92	8.97	11.96	4.20	16.45	13.66	80.4	9 6	5.00	80.0 20.0	16.05	9.92	5.06
4.16	18.23	7.07	1.2.2	79.1	18.13	0.57	0.97	87	7.06	06.7	60.0	2.03	14.7	0.21 0.59	2.69	1.23	0.83	1.27	3.02	5.64	0.35	1.25	3.13	12.26	1.87	130	. c	00.0	 	60.7	, 6	67.7	6.83	11.07	4.52	1.57	7.23
7.89	787	6.7	7.83	7.78	7.74	7.73	27.7	) ; r	0,7	69'2	7.67	7.67	7.66	7.65	7.64	7.64	7.61	7.60	7.57	7.54	7.53	7.49	2.78	7.46	, t		Į ;	14.7	7.36	7.35	7.34	7.31	7.29	77.7	7.27	7,24	7.23
27.0	3 6	PS./	5.20	21.45	1.06	710		1.00	2.05	0.08	22.51	0.53	0.25	<del>2</del> .	23.26	18.06	90.0	5.46	12.06	0.08	7.28	( <del>*</del>	2 7	17:6	0.0	2.44	6.14	0.31	8.77	4.	9.91	8.19	28.27	15.11	3.80	0.35	2.20
	9. ;	9.94	99.0	2.76	0.14		70.0	0.24	0.27	0.0	2.93	0.07	0.03	0.13	3.0g	2.36	0.01	0.72	1.69	, c	700		9 9	0.70	0.01	0.33	0.83	<u>8</u>	1.19	09.0	1.35	1.12	3.88	2.08	0.52	0.05	0.30
,	N62132	AA826324	R50775	W74362	144040	21211	AA156795	AA453437	H57082	AA463982	AA700739	H10231	H10068	AA780365	AA088886	W72666	195804	AA291773	56497	100000	AA/02602	K53455	AA633835	AA708441	AA485749	AA702973	R12473	R49644	AA521384	AA011390	N62936	AA459008	T66816	AA706968	R42946	N22766	2770
	287598	1420830	38833	245090	06687	200	502444	788217	204644	810661	435351	46927	46740	1035664	561971	345743	120695	725340	14780	20014	448032	39833	868169	505018	811166	447167	128243	37538	826995	429519	278729	814288	66317	451907	31299	26631	
	<b>6</b> 6	695	969	507	9	260	669	8	701	702	503	8	705	106	707	708	9 2	3 5	2 ;	E :	712	73	714	715	716	717	718	719	720	721	Ħ	773	5 2	7 2	3 5	9 5	77

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

	8	2	7	~	7	7	7	7	7	7	7	8	7	7	7	7	74	7	7	64	8	7	7	7	7	8	7	7	7	8	8	2	7	74	8	
1	6.82	6.11	6.43	11.27	15.36	16.05	14.61	10.38	11.80	7.84	7.62	5.32	8.84	0.77	6.81	0.27	4.28	11.87	14.49	3.14	9.27	10.38	15.45	1.16	8.65	12.53	8.19	13.46	8.34	11.61	3.78	5.09	6.48	15.52	3.19	
ar cell lines.	13.06	12.65	13.46	7.61	3.28	3.21	4.52	8.83	8.75	0.03	11.28	2.34	11.35	15.66	12.50	17.83	14.80	7.50	5.67	16.93	9.22	8.38	1.38	5.14	9.72	1.10	1.51	2.38	8.48	7.15	2.82	13.88	12.71	3.80	16.50	
2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines	1.81	2.92	1.78	2.72	2.94	2.29	2.42	2.25	0.87	13.51	2.47	13.68	1.14	4.89	2.00	3.04	1.98	1.68	0.85	0.86	2.37	2.09	3.97	14.50	2.34	7.04	10.95	4.74	2.73	1.71	13.90	1.52	1.22	1.08	0.68	
gher in at lea	7.23	7.23	7.22	7.20	7.19	7.18	7.18	7.17	7.14	7.13	7.12	7.11	7.11	7.11	7.11	7.05	7.02	7.02	7.00	6.98	6.95	6.95	6.94	6.93	6.90	6.89	5.88	5.86	6.85	6.84	6.83	6.83	6.80	6.80	6.79	8
10 fold or hi	1.17	8.99	0.31	4.64	11.27	8.56	0.24	15.96	0.11	105.24	14,44	0.62	1.51	3,14	95.95	0.26	2.95	2.01	0.28	1.75	0.68	36.26	0.05	0.07	3.78	90.0	2.74	23.86	4.75	0.83	3.35	0.25	2.79	4.70	5.63	
2 of 3 or	0.16	1.24	9.0	9.64	1.57	1.19	0.03	223	0.02	14.77	2.03	60.0	0.21	9,44	13.50	0.04	0.42	0.29	9. 5.	0.25	0.10	5.22	0.01	0.01	0.55	0.01	0.40	3.48	0.69	0.12	0.49	25.00	0.41	0.69	0.83	
٠	H19234	AA135809	AA011182	AA45521	AA431773	AA131909	AA128005	AA504201	AA677655	AA894557	AA455237	AA449118	R96520	AA455087	N66158	AA029722	AA598849	N53940	W72556	AA417288	T68878	AA485453	AA486741	<b>T58146</b>	AA886739	AA689222	AA452278	T63980	AA129135	AI025259	AA935560	AA398406	H56903	R23246	AA455940	
			359722	809828	782503	504308	501868	825282	897271	1416782	814779	785845	199644	812294	278531	366778	898333	247050	345330	731139	82215	811067	841221	79254	1500162	856585	787938	79761	586650	1641367	1556526	726791	204580	131094	813275	
	729	33	Ę	732	733	ሂ	735	736	737	<b>5</b> 5	739	740	741	742	743	74	745	748	747	748	749	750	751	752	753	754	755	756	757	758	759	£	761	79,	8	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

7	7	7	73	2	7	7	7	2	. ~	7	7	2	73	7	2	7	7	7	2	7	2	7	7	~	7	7	2	2	7	7	2	2	2	7		
1.76	0.98	8.83	3.12	7.33	122	13.94	14.72	4.64	4.14	4.12	8.09	1.85	9.54	4.75.4 4.00.4	4.81	1.28	9.68	6.74	13.83	5.55	6.04	3.84	2.39	12.67	7.59	8.78	6.08	9.70	1.28	12.22	4.18	8.37	3.99	12.04		
15.10	7.25	1.82	16.41	10,73	14.81	4.25	3.97	14.02	13.65	1.82	0.58	10.05	9.16	12.93	13.90	13.84	7.59	41.73	4 25	11.59	2.65	13.27	3.01	18.	9.29	9.13	11,14	6.70	12.25	8.08	1.59	7.88	13.18	5.53		
¥7 6	7 6	12.00 9 56	9 5	, c	3 +	. 87	5.5	3 4	3	177	12.51	7 07	1.31	c	97.7	1.02	0.4	\$ ;	74.	£	2.33	2.5	7 7 7	50.7	9 5	2.30 8.50	50.5 20.5	2.85	69.4	666	0.50 13.18	7 6 7	2.67 4.05	S 25		
,	6.77	6.77	6.74	6.73	0.7.0	9.7	D (0)	0 0 0 0	0.07	5.67	6.65	6.0 40.0	6.62	0.07	6.58	6.58	6.58	6.57	6.51	6.51	6.50	6.49 9 . 0	6.49	ò .	0.4/	5. 5.	9.45 5.45	0.42	14.0	6.41	6.39	6.37	6.37	6.5 2.4	<u> </u>	3
	7.09	1.65	10.08	9.88	7.44	3.87	421.08	3.42	3.96	3.02	17.94	7.75	0.07	63.60	2.64	1.48	1.57	0.12	0.24	5.27	6,55	8.71	10.14	2.76	0.07	0.12	27.63	0.45	6.74	0.21	1.66	98.6	3.76	0.1	4 X	
	1.05	0.24	1.50	1.47	Ξ	0.58	62,96	0.51	0.59	0.45	2.70	1.17	0.01	9.61	0.40	0.22	0.24	0.02	<b>0</b> .04	0.81	1.01	¥.	1.56	0,43	0.0	0.02	4.30	0.07	1.05	0.03	0.26	1,55	0.59	0.02	7.0	
	N48707	AA454562	AA454597	N47008	AA398949	H05085	AA857413	R53446	W37447	AA883391	AA165678	AA424790	AA448400	155728	AA142869	AA029997	H09997	AA039851	151617	N79745	R43328	AA953357	AA598796	R32406	AA129677	AA428186	AA628788	AA452542	AA448664	H15910	R89846	N25938	N39611	AA485351	AA405533	
	279397	809503	_		-	43759		40038	321900	1461316										300276	32134	1573251	860868	135688	565493	773564	877784	788488	786078	159462	194717	268711	277.172	810996	772410	-
	2	99	99	29	89	69	2	7	22	5	74	75	9/	E	178	677	280	781	782	783	8	785	786	787	. 887	789	790	791	792	793	794	795	796	797	798	

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

2	~	2	7	7	7	2	7	7	~	2	8	~	8	~	~	7	~	8	7	7	7	~	~	~	7	7	7	~	~	~	~	8	2	~	
6,13	11.23	1.13	11.81	9.86	6.32	8.17	7.26	8.95	7.88	10.08	13.49	5.30	1.69	5.48	9.06	9.66	3.69	6.77	2.74	6.80	7.78	10.29	9,56	9.04	3.85	1.49	7.48	7.07	11.91	5.03	1,00	12.34	12.62	4.79	
12.20	6.38	4.43	4.38	8.03	10.78	8.05	10.13	2.46	8.82	6.21	5.01	0.27	3.53	12.34	8.92	2.60	12.42	0.76	3.72	10.69	2.18	7.55	2.69	7.76	13.05	12.18	9.73	8.86	3.64	11.73	3.46	4.07	0.62	11.45	
0.76	1.48	13.51	2.86	1.06	1.79	2.64	1.45	7.40	2.11	2.51	0.27	13.18	13.52	0.87	0.68	1.35	2.28	11.05	12.11	1.20	8.48	0.57	6.15	1.59	1.47	4.68	1.12	2.39	2.76	1.52	13.72	1.70	4.85	£.	
6.36	6.36	6.36	6.35	6.31	6.30	6.29	6.28	6.27	6.27	6.27	6.25	6.25	6.24	6.23	6.22	6.20	6.20	6.20	6.19	6.16	6.15	6.13	6.13	6.13	6.12	6.12	6.11	6.11	6.10	6.09	90.9	6.03	6.03	6.02	22
1,50	6.46	3.38	26.80	4.87	51.96	5.87	2.21	1.37	1.58	0.15	2.29	13.69	4.98	4.11	1.36	12.01	3.99	0.08	8.84	0.33	2.57	15.94	12.43	2.89	10.01	0.15	2.40	16.02	207.98	26.54	90.0	0.70	5.33	16.34	
0.24	1.01	0.53	4.22	0.77	8.25	0.93	0.35	0.22	0.25	0.02	0.37	2.19	0.80	0.66	0.22	1.94	0.64	0.01	1.43	0.05	0.42	2.60	2.03	0.47	1,63	0.02	0.39	2.62	<b>%</b> .08	4.36	0.01	0.12	0.88	271	
R93068	AA465223	AA777555	171991	AA459244	AA042812	W85878	N63988	N77828	H54659	R44201	N24869	AA430367	AA877997	AA701978	R38539	H22568	AA630373	W31389	AA779949	N56875	AA865924	AA630104	AA430744	AA609891	AA608752	R26456	R92186	AA430052	AA410517	AA459106	R44617	AA775557	AA165313	AA453441	
196824	814211		85409	81444	486538	416075	289496	302221	203179	34254	267135	769857	. 1475308	435957	23073	52128	854897	320602	461933	277476	1470195	854701	770992	1032080	950945	133192	195817	781109	753862	814303	33342	378016	594693	788225	
199	908	8	803	803	Ş	808	808	807	808	608	810	811	812	813	814	815	818	817	818	8	820	821	822	823	824	825	826	827	828	829	830	<b>8</b>	832	833	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

14 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	0.39 6.02 2.14  7.71 6.00 2.89  0.54 5.99 4.63  8.34 5.98 2.44  2.39 5.98 2.44  2.39 5.98 2.42  0.49 5.98 2.42  0.49 5.98 2.42  0.55 5.96 2.30  0.55 5.96 2.30  0.50 5.93 0.97  1.85 5.91 1.63  1.85 5.93 11.41  1.85 5.93 11.41  1.85 5.93 0.97  0.36 5.89 2.20  0.36 5.89 0.84  0.36 5.89 0.84  0.36 5.89 0.84  0.41 5.89 0.84  0.41 5.89 0.30  0.41 5.89 0.30  0.41 5.79 0.30  0.42 5.79 1.15  0.59 5.79 1.15  0.60 5.80 2.24  0.70 5.83 3.05  0.70 5.83 3.05  0.70 5.79 1.15  0.70 5.79 0.30  0.70 5.79 1.15  0.70 5.79 0.30  0.70 5.79 1.10  0.70 5.79 1.10  0.70 5.79 1.10  0.70 5.79 1.10	6.02 6.02 6.03 6.04 6.09
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Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

8	8	2	8	2	7	7	7	2	8	7	7	8	7	<b>~</b>	7	2	7	8	7	2	8	7	8	8	2	7	7	8	2	. ~	8	~	7	8	
10.01	3.07	5.38	1.55	9.54	1.87	3.16	7.73	11.69	12.25	0.46	6.19	5.00	8.55	11.39	2.59	3.58	7.48	5.99	5.46	2.47	4.42	4.36	7.67	0.55	8.52	6.45	3.69	8.81	7.85	1.70	12.10	7.61	3.66	10.42	
6.27	12.44	9.62	9.78	6.51	9.35	12.71	8.12	4.30	1.00	12.66	2.36	1.37	7.77	0.44	8.42	11.49	1.28	10.00	9.88	7,41	11.14	1.27	2.85	12.83	7.04	8.01	10.54	60.0	7.74	7.64	6.	8.14	10.90	1.50	
0.89	1.62	2.14	5.72	1.00	5.83	1.17	1.17	1.02	3.76	3.85	8.40	10.55	0.55	5.02	5.85	1.75	8.06	0.80	1.43	6.89	1.18	11.02	6.12	3.23	1.02	2.10	2.27	7.57	0.87	7.11	3.34	0.67	1.85	4.45	
5.72	5.71	5.71	5.68	5.68	5.68	5.68	5.68	5.67	5.67	5.66	5.65	5.64	5.62	5.62	5.62	5.61	5.60	5.60	5.59	5.59	5.58	5.55	5.55	5.53	5.53	5.52	5.50	5.49	5.49	5.48	5.48	5.47	5.47	5.46	56
1.35	1.90	0.15	1.58	5.71	2.92	7.88	2.35	7.19	90.0	0.81	18.53	6.68	0.25	0.10	90.0	0.18	17.75	0.07	0.17	4.03	11.01	3.89	3.65	0.05	132.57	0.41	5.42	66.94	11.11	8.93	0.05	5.38	7.03	2.28	
0.24	0.33	003	0.28	1.00	0.51	1.39	0.41	1.27	0.01	0.14	3.28	1.19	0.05	0.02	0.01	0.03	3.17	0.01	9.03	22.0	1.97	0.70	99.0	0.01	23.98	0.07	66'0	12.19	2.02	1.63	0.01	9.08	1.29	0.41	
H37817	R60223	R40481	R45952	N72274	N51498	AA131707	R89363	AA775259	AA011389	R51382	AA699560	AA669218	AA757588	AA427400	W93500	AA679907	AA608730	W96197	H47315	T98201	AA452809	H05563	AA035580	AA443286	AA670408	AA907714	R22206	AA487608	R22308				AA504262		
191546	42842	27817	34063	291374	282104	503865	196037	878600	429517	39219	433474	856592	395609	770979	357344	869375	920036	361688	193139	121798	789049	43977	471707	783987	878788	1505898	130858	839081	130845	241432	179556	197206	825408	525799	
869	870	178	872	873	874	875	876	877	878	879	880	881	882	883	884	885	886	887	888	883	830	891	892	893	894	895	896	897	868	668	006	901	905	803	

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

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•	<b>4</b> (	7	7	8	7	7	2	7	2	7	7	63	7	7	7	7	8	7	7	8	7	7	7	7	7	8	7	7	7	7	7	7	7	7	7		
	6.30	7.49	8.44	5.51	8.69	17.7	11.81	3.53	4.35	8.77	4.15	11.07	8.24	4.21	6.03	7.16	7.71	10.03	8.09	5.20	7.18	5.27	6.95	10.80	7.50	7.02	11.04	8.43	6.43	7.68	0.55	1.88	3.74	1.61	6.64		
	8.29	6.97	6.79	8.15	5.96	6.69	3.31	10.54	10.79	5.32	10.96	1.00	6.68	10.44	1.01	7.63	7.20	3.73	5.86	9.50	6.76	8.00	868	0.18	2.95	2.97	4.37	5.07	7.32	5.49	10.92	10.57	1.80	10.71	5.99		
	1.75	1.86	1.09	2 62	19	2 2	5 5	2				507	3 5	2		 	<u> </u>	30:- 20:-	ž 70 c	, t	5. 5	 	3 6	<b>1</b> 8	, Y	5 K	9 6	5 6	5 5		, c	07.4	2 <b>5</b>	2.27	2.85	<u>.</u>	
•	5.45	77.7	77	ţ <u>Ş</u>	, r	24.0	4	0.40	 	بن رو پی در	0 0	υ, r ο α	5.50			5.32	5.31	5.31	5.30	5.30	5.29	5.29	67.0	5.28	2.5	5.20	9.40	5.20	97.0	5.23	57.5	5.22	5.21	5.20	02.6	<u>.</u>	27
	0.26	) i	6.03	1.05	0.45	33.51	0.69	<del>.</del> 6	0.12	2.76	7.13	11.29	0.05	1.61	0.28	0.05	6.28	2.11	14.65	21.66	4.03	0.23	20.12	0.41	4.49	3.68	41.0	0.15	0.58	44.95	15.25	0.08	1.87	0.76	0.46	0.35	
	4	60.0	0.01	0.19	0.08	6.18	0.11	0.18	0.05	0.51	1.32	2.10	0.01	2.17	0.05	0.01	1,18	0.40	2.76	4.09	0.76	9.0	3.81	90.0	0.85	0.70	0.03	0.03	0.1	8.60	2.82	0.02	0.36	0,15	0.09	0.07	
	,	AA428514	H96356	R89287	4A884428	AA706829	AA883100	AA969184	173031	H18630	AA458827	AA497034	R44292	AA402965	W72310	R73525	R16676	AA598668	AA775865	AA455652	N21334	N57858	177840	AA628190	W72679	H59305	AA879474	N20338	AA620614	AA133166	AA169645	AA404290	AA284280	W32272	AA121504	AA446748	
				195786	1466998 A	451805 A	1467751 A							741790			129664					247089						264646	1048685	490753	594517	758360	324665	321386	489881	784128	
		z	8	8	20	80	8	5	=	7	ū	7	5	16	11	8	916	320	52	22	323	924	925	926	827	928	929	930	931	932	933	934	935	936	937	938	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

2	7	. 4	~	7	. 14	. 24	. 64	. 6	2	. 4	~	7	~	7	7	٠ 7	~	~	7	8	~	8	7	7	8	2	. 2	. 7	7	. 7	. 7	7	7	٠ ٦	I
6.90	5.15	7.33	10.09	7.94	10.01	3.58	8.34	5.28	5.12	7.63	0.32	5.48	5.67	7.92	5.79	1.87	3.42	6.61	5.79	0.97	5.65	6.75	6.79	2.27	6.33	6.25	5.46	7.16	10.79	6.59	1.00	5.32	1.00	6.60	
6.91	9.14	1,00	1.15	0.80	3.54	11.08	5.30	8.40	9.02	5.29	8.82	8.12	8.65	5.51	8.28	6.10	10.25	6.33	8.07	5.74	8.14	6.08	6.72	7.27	7.99	6.45	7.62	5.39	3.68	5.47	5.04	8.39	5.50	5.68	
1.72	1.23	7.12	421	6.70	1.82	0.72	1.68	1.62	1.14	2.35	8.08	1.54	0.82	1.63	0.91	7.00	1.25	1.99	0.98	8.10	0.97	1.91	1.20	5.15	0.37	1.97	1.57	2.09	0.13	2.45	8.44	0.71	7.91	2.12	
5.18	5.17	5.15	5.15	5.15	5.12	5.11	5.11	5.10	6.09	5.09	5.07	5.05	5.04	5.02	4.99	4.99	4.97	4.97	4.95	4.94	4.92	4.91	4.90	4.90	4.90	4.89	4.88	4.88	4.87	4.84	4.83	4.80	4.80	4.80	28
9.47	0.51	0.05	22.27	90.0	9.93	0.07	0.19	0.43	32.60	1.98	16.55	40.76	770	3.45	4.54	8.53	26.53	1.93	6.34	0.05	8.75	0.30	2.11	90.0	8.32	0.71	1.49	34.71	28.69	10.43	9.05	13.99	0.05	92.90	
1.83	0.10	0.01	4.32	0.0	<del>2</del> .	0.01	9.0	90.0	6.40	0.39	3.26	8.07	<b>6</b> .0	0.69	0.91	1.7.1	5.33	0.39	1.28	0.01	1.78	90.0	0.43	0.02	1.70	0.14	0.30	7.11	5.90	2.16	0.01	2.91	0.01	19.35	
AA644334	AA426561	AA398218	AA129931	W84774	H99997	R38169	AA683338	AA777779	AA148505	R71689	AA962541	AA421230	AA490605	N49899	H61552	AA149637	AA504631	N20480	R44346	H39560	AA478542	R82041	N56982	4A114106	H16743	AA904969	R55750	A035384	A043228	N62372	VA454554	W064627	R43271	AA456931	
-	_	•	••	Ī	263342	23658	1293016	448920	491524	155575	1602209	739247	824109	243653		_	_						277545		50114	•				_	_	_		838568 A	
88	95	<u>\$</u>	942	943	\$	945	946	94	<b>8</b>	949	820	951	852	953	954	955	926	957	928	828	960	961	962	963	96	865	998	2967	898	696	926	971	272	£	

Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prostate

2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.		-								
87.4	258966	N31641	1.57	7.55	4.80	48.	5.26	7.20	7	
1 10	503715	A4131578	5.14	24.62	4.79	2.51	6.24	5.62	7	
6.76	14504B	R72434	0.02	0.10	4.78	3.62	10.08	0.65	~	
	464479	DEABAR	0.72	3,44	4.77	1.60	6.79	5.93	7	
	204430	DE STATE OF	1 6	99	4.75	1.18	6.82	6.25	7	
<b>20</b> C	625089	A553413	0.13	09'0	4,75	1.30	6.47	6.47	7	
D (	600070	1212000	200	4.36	4.75	2.49	5.38	6.37	7	
980	652521	H002/5	3.47	£ 44	4.74	5.66	0.08	8.48	2	
981	453005	AA//9165	÷ ;	, F.	4.74	2.27	5.74	6.21	7	
382	50000	AA/00419	3 6	7 27	4 72	2.52	5.51	6.12	7	
983	43826	H05/68	0 6	ţ ;	467	6.88	1.33	5.80	8	
<b>8</b> 8	432732	AA701587	3 6	 	99.	6.62	5.90	<u>+</u>	7	
985	882488	AA676590	9 6	7 6	20.	78	2.63	5.38	7	
986	46097	H08796	0.39	1.62	3 5		5.33	6.39	7	
282	809466	AA443094	0.90	4.18	30.4	61.7	5.63	88	7	
988	745192	AA626847	1.12	5.17	7.07	2.30	8 6	A.C. A.	•	
686	898083	AA598797	3.37	15.49	4.59	2.32	DZ (	0.43	• •	
066	796606	AA460521	0.97	4.45	4.59	2.55	5.24		<b>v</b> (	
166	327245	AA284291	0.98	4.50	4.59	2.99	5.72	5.05 1	7 (	
90	744044	AA629251	0.01	0.06	4.58	5.99	0.75	7.00	7	
1 6	882355	AA676298	2.64	12.03	4.56	6.34	5.06	2.28	2	
	2615R7	HORGSS	2.99	13.59	4.54	1.15	6.62	<b>5</b> .85	7	
, y	R24728	AA488979	=	5.05	4.54	0.93	7.47	5.23	7	
200	283208	N51367	0.59	2.67	4.54	0.99	7.39	5.23	7	
999	453220	AAZDABSB	0.08	0.38	4. 2.	1.08	5.53	7.00	7	
	112056	B24451	0.65	2.96	4.53	2.21	5.84	5.54	2	
000	731034	AA421270	8,95	40.40	4.52	2.12	6.30	5.12	7	
666	1468466	AA885096	0.60	2.71	4.50	5.38	2.55	5.56	2	
3 5	206277	H58542	1.42	6.39	4.50	1.27	5.28	6.85	7	
3 8	75650	T58434	10.1	4.53	4.49	2.28	6.17	5.02	7	
	257312	N29624	0.36	1.60	4.48	0.90	6.98	5.57	7	
3 8	1048899	AA778562	0.07	0.32	4.48	1.56	5.09	6.78	7	
Š	51700	H22854	0.24	1.09	4.47	6.45	5.25	1.71	<b>7</b>	
900	1466599	AA883660	0.12	0.53	4.46	1.21	6.24	5.83	7	
200	51328	H20743	0.01	90'0	4.4	2.31	5.02	5.98	7	
800	1504005	AA904776	0.11	0.49	4.	1.72	5.71	5.87	2	
3					29					
					ì					

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

5.86 0.44 6.47 5.84 5.62 5.79 5.36 5.76 6.87 5.58 5.36 5.41 5.49 5.65 6.01 5.93 5.55 5.01 0.56 5.58 5.46 3.55 5.01 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines. 5.14 6.04 6.06 6.96 6.96 5.51 5.51 5.77 5.61 5.44 5.55 1.07 5.50 5.73 1.63 6.05 5.14 5.12 1.78 5.82 5.04 5.12 2.19 6.57 2.44 2.38 1.74 1.92 5.57 0.99 1.50 45 1.28 1.08 1.91 1.25 5.58 1.40 1.38 5.38 1.58 8. 0.57 1.23 0.97 5.88 5.25 5.25 1.13 1.12 1.12 1.01 0.43 4.32 4.32 4.32 4.28 4.08 4.28 4.25 4.24 4.17 4.13 22.74 11.37 8.82 5.73 11.43 0.40 0.42 0.10 0.39 0.27 6.07 1.69 9.76 0.57 9.78 0.36 0.83 4.55 4.25 7.30 0.07 0.02 5.27 0.09 0.06 1.43 2.68 2.10 1.37 2.75 0.03 0.41 2.36 0.14 2.38 0.20 1.12 3. 3. 0.02 2.77 AA205320 T70122 **4A398355** AA404338 AA112979 AA884382 AA626247 AA625791 AA443300 AA437245 AA708431 AA487206 AA633993 AA862814 AA485692 AA883755 A676598 H08227 N50828 N35579 R60949 N50935 R07012 A045340 T90621 N79230 T54527 T62552 H79566 H84915 N58136 H47089 N32071 1461715 530545 758302 745523 784589 466824 744911 647816 757463 814906 281103 110772 80946 280309 272238 42400 505997 841287 126695 858292 1469211 45514 73659 79726 292388 382483 249517 247587 93420 239661 187141 012 1013 1014 1015 1016 1018 1025 1026 1027 1028 503 1017 019 92 1021 1022 1023 1024 629 630 631 632 £ 5 037

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	-	<b>-</b>	-	<del>-</del>	-	<b>-</b>	-	<b>-</b>	-	-	-	-	-	-	-	<b>,.</b>	-	-	-	-	<b>-</b> -	-	-	-	-	<b>-</b>	-	-	-	-	<b>-</b>	<b>-</b>	-	-	-	
	1.00	0.59	1.33	1.09	1.00	2.22	2.96	2.07	0.74	2.83	1.09	1.04	0.74	0.30	2.47	1.77	0.29	0.76	0.38	2.	1.00	0.97	1.10	0.36	2.35	1.26	2.37	0.68	0.69	0.07	0.48	1.06	1.96	1.21	0.19	
lines.	1675.21	1262.88	1231.62	761.20	730.07	708.49	652 FB	586.88	515.39	458.08	446.45	438.73	264.42	241.37	221.03	20 02	218 92	218.05	214.82	212.83	191.25	176.58	175.32	173.57	168.03	162.70	161.00	153.57	149.67	151.46	148.64	143.86	135.65	135.09	137.23	
2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	7	17.1	800	25.5	n 4 - c	2.70	2.72	2.43	2.91	6.33	20.1	1.13	2 7	47.1	600	B	P) 1	0.7	0.60	2.48 7.0	2.70		5 4		3 5	1.13	0.86	28.0	2.74	50	88.	9.0	0.62 0.62	60.7	5. co	D S
er in at least 1	6	559.26	421.73	17.114	254.59	244.61	237.81	219.32	197.29	172.23	154.17	149.45	146.97	88.80	80.84	75.00	74.54	73.32	73.13	72.56	72.19	64.60	59.54 40.65	27.65	28.52 28.53	54.79 E0.25	20:00	7. 7.	2. E	20.00	50.64	50.20	48.61	46.77	46.43	48.12
) fold or highe		5.59	7.11	158.35	2.55	2.45	2.38	2.19	1.97	6.87	6.56	73.56	49.37	69.16	7.42	3.61	8.93	16.59	23.96	1.43	2.83	0.65	3.72	20.73	18.15	1.84	14.22	3.65	4.65	0.94	7.71	1.05	9.24	32.83	1.25	12.55
2 of 3 or 10		0.01	0.02	0.39	0.01	0.01	0.04	0.01	0.0	9. 20.	9.0	0.49	9.3 24.	0.78	90.0	9.09	0.12	0.23	0.33	0.02	9.0	0.01	90.0	0.35	0.31	0.03	0.26	0.07	0.03	0.02	0.15	0.05	0.19	0.70	0.03	0.27
re Cancer Cen Lines vs. Nor.		N47961	N93470	N48913	N30131	R01937	W20486	AA476502	AA486538	AA875888	AA432058	AA456130	AA406070	AA401342	AA461320	AA426212	H11642	AA598983	N33323	AA399245	H05961	R61395	AA133554	AA002166	AA171784	H16537	AA432278	AA459400	AA481806	AA001219	AA621324	H47076	AA013353	AA398384	AA443140	AA456951
		280291	307328	279613	268115	124567	327495								796334	769542	47963	897722	270603	726439	44007	37986	586845	427693	594683	49131	782353	810959	838877	362278	1048794	193087	360355	726934	798723	838600
ie Caŭ		1044	1045	1046	1047	5 2	3 2	1050	193	1052	1053	1054	1055	1056	1057	1058	1059	900	1061	1052	1063	1064	1065	1066	1067	1068	1069	1070	1071	1072	1073	1074	1075	1076	107	1078

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

108.31 2.03 1.83 9.84 0.39 0.18 1.65 1.65 95.10 2.06 1.78 1.00 2.37 **0**.86 0.39 1.00 0.25 2.69 0.76 1.56 128.29 129.35 122.05 122.87 120.12 117.01 117.08 14.00 113.82 1.33 1.22 110.55 109.30 106.62 0.03 106.86 104.16 1.02 99.56 98.13 8. 90.43 92.44 90.70 91.71 84.23 118.16 118.37 1.18 1.76 0.85 1.15 0.97 0.73 0.68 0.12 1.05 1.08 1.73 2.92 43.69 41.59 41.07 40.88 40.15 40.03 39.71 39.20 39.51 37.26 36.69 38.61 36.82 36.32 36.21 35.93 8.13 33.51 32.37 31.55 31.48 31.48 31.06 28.99 28.51 15.59 23.11 14.33 10.92 30.27 82.43 2.07 å Ş 1.73 13.35 16.34 10.68 13.40 1.94 0.34 0.32 0.32 37.35 0.31 1.21 5.31 0.36 0.28 0.78 2.21 0.06 0.45 0.05 0.30 0.39 0.0 0.0 1.18 0.01 AA425743 AA180453 AA481788 N34482 W187595 AA677643 AA457138 W86779 H08850 W4299B N64840 W88720 N26083 N27641 H23524 N34943 AA449300 AA704816 R10140 N23865 N46845 N52158 H57136 AA159497 AA128407 AA121387 AA872602 N48271 W43028 N66454 R37289 625516 416808 255659 268979 323251 284701 46055 417760 51799 609438 838853 276905 897262 810459 128461 254545 284408 279176 204686 785866 1323328 282161 592403 526567 452848 329059 323448 564231 285581 773678 1084 1085 1086 080 1087 1088 1089 1092 1091 1093 <u>1</u>09 1095 1098 5 1096 6601 101 102 豆 호

Table 2-10; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

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A C 34 A 4	2 of 3 or 10 fold or higher in at least 1 of 3 calluct certainer.
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	or higher
	r 10 fold
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2.63	1.55	2.80	1.17	0.80	1,68	2.46	; c	00.0	2.30	7.4.7	1.00	2.23	2.10	3. t	7.1	7.7	2.11	1.45	0.77	0.10	0.87	0.97	2.57	1.38	1,40	0.63	0.68	1.70	59.09	0.12	1.09	0.84	2.16	0.46	0.59		
75.29	73.65	73.28	74.73	74.38	22.44	70.67	70.01	טר.ר/	71.45	71.41	70.65	71.20	68.08	69.82	88.24	67.64	64.57	65.48	66.20	65.41	64.81	64.73	63.34	64.28	62.07	63.23	62.26	59.72	2.08	60.67	58.64	59.31	56.55	57.49	57.17		
0.1	2.60	1.21	0.75	200		<u>ş</u> ş	1.92	- -38	0.41	1.19	2:32	820	2.54	<del>1</del> .0	1.52	128	1.35	1.39	0.83	1.64	1.16	0.97	0.70	0.84	1.59	1.18	0.56	12.	1.29	1.10	1.33	0.81	0.96	4.	1.24		
26.31	25 93	25.76	2 3	25.33	53.10	25.13	25.02	24.82	24.72	24.67	24.66	24.57	24.27	23.82	23.68	23.36	22.90	22.78	22.60	22.38	22.28	22.23	22.20	22.17	21.69	21.68	21.17	20.87	20.81	20.63	20.35	20.32	19.89	19 79	19.67	33	;
26.72	23.43	35.40	0.20	90.00 100.00	15.3	9.24	4.63	5.49	59.39	. 16.06	0.25	0.84	10.63	37.42	10.35	12.38	27.72	6.49	9.38	2.19	14.30	17.18	0.81	8.49	4.05	12.75	29.89	10.68	0.18	1.66	3.00	5.45	20.5	12.57	1.90		
5		<u> </u>	io 1	0.15	60.0	0.37	0.18	0.22	2.40	9.65	0.01	0.03	4	1.57	0. 44	0.53	177	0.28	0.42	0.10	200	0.77	8	0.38	0.19	0.59	1,42	0.51	0.0	0.08	0.15	720	2, 0	3 6	5 6		
,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	AAK / 1004 I	N74889	R40057	AA418028	AA425214	AA417280	H15050	R49462	H97508	N67571	N64774	AA432096	44884321	R09504	AA011570	AA022668	A 4 2 5 0 3 0	AA417211	AA490264	R92512	AA406220	MERAGE	MAKANI	SOUTH	W74696	AA470073	H89563	N62079	AA138049	R49436	010478	202020	739304 e	AA885141	LC862N	78182	
	080/6	299603	27544	767739	773293	731115	48363	38260	251961	285682	284569	784154	1455803	127711	429660	364454	757735	731154	923705	106475	242800	VOCAT'	200002 208745	2000	244747	730377	253534	289865	50203	18744	32046	1000	10010	1466531	206802	- -	
:	1114	1115	1116	1117	1118	1119	1120	4121	1122	1 5	2 2	175	2 2	1120	128	2 5	1 1 1	130	5 5	7611	555	5 5	1155	0011	127	130		144		7	3 3	\$ :	1145	1146	1147	**	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

	<del>-</del>	-	-	-	-		-	-	-	-	-	-	-	-	-	-	-	-	<b>-</b>	-	-	-	<b>-</b>	-	-	-	-	<b>-</b> -	-	-	-	-	-	-	-	
•	8.	1.97	54.90	0.71	0.76	0.38	0.41	<del>.</del> 2.	1.4	1.23	1.83	0.50	0.53	1.45	1.64	2.29	0.63	2.70	0.69	1.59	2.06	1.55	0.77	0.74	1.16	0.53	1.30	2.52	44.37	2.06	1.23	1.07	0.38	0.79	78.0	
	55.61	53.79	1.00	53.32	54.30	54.14	53.43	51.75	51.86	1.71	49.38	49.70	50.90	0.87	48.99	46.96	49.51	1.50	0.05	46.45	45.77	46.79	47.42	46.41	46.01	46.29	44.76	42.94	1.00	42.61	43.72	42.03	44.40	43.90	42.90	
	1.00	1.56	1.10	2.52	0.98	0.36	0.31	1.27	0.55	50.03	1.35	2.13	0.40	49.43	0.80	1.89	0.36	45.76	48.82	1.51	1.72	0.92	1.01	1.38	1.23	1.38	0.93	1.31	1.32	1.79	1.18	2.79	1.07	1.08	1.48	
	19.20	19.11	19.00	18.85	18.68	18.29	18.05	18.02	17.94	17.66	17.62	17.44	17.28	17.25	17.14	17.05	16.83	16.65	16.52	16.52	18.51	16.42	16.40	16.18	16.13	16.07	15.67	15.59	15,56	15.49	15.37	15.30	15.28	15.26	15.12	용
	0.19	8.40	0.19	0.19	12.10	0.48	14.96	2:32	17.51	7.11	6.75	1.37	3.05	8.13	4. 48.	6.13	4.60	5.82	3.57	6.03	5.21	3.70	11.22	4.98	14.63	1.63	4.23	5.43	0.16	2.75	8.40	7.61	3.82	8.56	12.92	
	0.01	0.44	0.01	0.01	0.65	0.03	0.83	0.13	0.98	0.40	0.39	0.08	0.18	0,47	0.11	0.36	0.10	0.35	0.22	0.37	0.32	0.23	0.68	0.31	0.91	0.10	0.27	9.36	0.01	0.18	0.55	0.50	0.25	0.56	0.85	
	-	AA167120		AA421256	T80832	R43189	AA884397	AA725397	H99075	•		AA188710				R60731	R43017	-	N50880	•	N51709	H98676	AA620301	R68409	N40202	AA190825	R51506	AA495938	N75473	A1005321	N66348	W44338	<b>VA845168</b>	VA421170	A083514	
	9 525478	9 609209	344618	731016	109275	32551	1461725	1342650	262060	743828	1048694	626186	745330	767449	451871	42225	31869	1468260	281003	731422	279102	261989	1030808	137853	276438	627401	38677	768445	299162	1626197	285370	323599	1412504 /	_	549157 A	•
	<del>+</del>	115	1151	1152	1153	1154	1155	. 1156	1157	1168	1159	1160	1161	1162	1163	1164	1165																1181	1182	1183	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

2000	A A B A A 17 A	0.33	5.05	15.11	0.85	43.54	 	
	A 449.25	2	7.34	14.83	1.04	42.32	1.13	
731444	A4412433	3 6	4.03	14.76	1.17	41.82	1.29	
30986	R41782	77.0	) i	14 65	1.26	42.02	0.68	
279616	N48302	0.58	0 1	20.7	133	40.40	1.92	
280270	N47954	0.53	5.7	0.4	2 40	41.08	0.42	
1474149	AA911971	0.02	0.34	40.4	) i	74 04	0.95	
80242	T64323	1.49	21.55	14.50	1.5	t : 0	14	
00000	A A 211 4 B 3	0.05	0.78	14.36	0.98	90.90	2 6	
000700	200000	5	1.87	14.30	1.45	41.37	90.0	
4503/5	AA/U3660	2 6	75.8	14.22	0.76	41.22	0.67	
784242	AA446882	6.0		43 00	2.13	1.37	38.45	
25933	R36969	0.01	0.14	000	91 +	39.26	1.48	
42433	R61189	0.15	2.05	9.60		0.11	0.89	
45501	H08397	0.09	1.31	13.90	50.0	. 6	53	
0,44,0	AAAETABT	0.58	8.10	13.87	0.91	39.10	2	
0 10	2010010	77.0	10.61	13.85	1.22	39.02	ري. ا	
284845	N66/24	3 6		13.84	2.64	2.34	36.52	
744417	AA621218	0.0	<u> </u>		1 10	39.01	1.27	
328689	W45330	0.70	9.05 40.0	5.5	100	39.43	0.85	
342089	W60894	- 80:	14.87	5.73		30.00	0.39	
38015	R61556	0.17	2.31	13.67	80.0	2 2	2.87	
810511	AA464541	<b>6</b> 0:	14.75	13.59	1.51	9.65	, F	
10000	100722	0.62	8.43	13.52	68.0	38.86	- (	
006707	133812	2	2.77	13.50	98.0	39.39	0.23	
549035	AAU83207		. 69 ^	13.50	0.89	38.18	1.42	
612782	AA181723	) ·	42.66	13.37	0.86	. 38.68	0.57	
730888	AA417031	0.65 0.65	60.21	12 37	1.42	38.10	0.57	
34901	R45114	0.27	3.60	5.5	;	38.03	0.79	
243410	N38992	0.90	12.03	13.35	4 6	28.24	0.65	
730398	AA469939	<u>2</u> .	13.46	13.30	0.80		2.48	
E07778		900	0.26	13.29	2.28	35.14	2 4	
13007	•	05.0	6.68	13.27	1.42	37.10	67.1	
7000	•	0	7.81	13.26	0.73	38.34	0.72	
283045			5 5	13.22	1.19	37.93	0.53	
796130	AA460967	<b>X</b>	S	1 4	0.42	38.93	0.11	
591814	AA143467	0.09	4.19	5.13		7 75	1.60	
30803	N92310	0.22	2.89	13.01	9/:	70.70	9 0	
		80	44.04	12.99	1.47	36.87	6.0	
4200	•	3	7.09	12.96	£.7	36.05	1.51	
1032072	2 AA609887	20.00	80.					

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

415820 358046 80186 454317 768111 299024 41300 278629 1465883 208413 782279 296448 511233 359040 366795 588159 742671 1031375 129862 813695 420584 203488 7456160 433350 112559 785788 346615 112559 785788 346615 785788 346615	1.62	0.98	0.65	1.30	1.58	1.39	1.00	1.26	2.43	0.71	0.76	0.79	35.74	1.66	0.95	1.50	1.46	1.28	0.64	32,33	1.25	0.58	0.80	1.06	1.61	1.68	0.95	1.66	0.40	0.27	1.51	0.90	1.13	. 0.32	•
415820         W84815         0.63         8.15         12.91           958046         W94691         0.52         6.66         12.84           454317         AA61822         1.07         13.65         12.72           454317         AA618968         0.67         8.45         12.72           768111         AA418968         0.67         8.45         12.60           299024         N75394         0.46         5.81         12.52           41300         R56642         0.21         2.68         12.40           278629         N86203         0.01         0.13         12.62           278629         N86203         0.24         6.06         12.40           278279         AA631750         0.52         6.39         12.47           28648         N74623         0.02         0.19         12.45           28649         AA402242         0.48         5.89         12.16           359040         W92278         0.48         5.89         12.16           358153         AA402242         0.48         5.89         12.16           358040         W92278         0.48         5.89         12.16           3581	36.14	36.21	36.50	1.48	35.01	35.31	35.57	35.21	33.25	35.13	1.49	34.77	0.66	34.01	34.90	33.59	32.82	33.72	34.33	2.58	33.75	33.81	32.40	32.62	32.51	31.51	0.28	1.2	32.53	33.21	30.58	31.68	29.73	0.83	25.43
415820         W84815         0.63         8.15           356046         W94591         0.52         6.66           80186         T64192         1.07         13.65           454317         AA677165         0.33         4.20           768111         AA418988         0.67         8.45           298024         N75394         0.46         5.81           510136         AA652239         0.01         0.13           41300         R56642         0.21         2.68           278628         N66203         0.49         6.06           1465883         AA684317         0.67         8.34           208413         H62162         0.32         0.96           278629         N664317         0.67         8.34           208413         H62162         0.32         0.96           29648         N74623         0.02         0.19           29648         N74623         0.02         0.19           511233         AA088701         0.18         2.21           359040         W92278         0.48         5.89           58159         AA135222         1.78         21.58           742071	0.97	<del>5.</del>	1.20	35.38	1.20	1.04	1.00	0.99	1.7	1.55	35.07	1.54	99.0	1.20	0.77	1.40	2.01	1.11	0.97	1.00	0.87	0.82	1.91	122	0.78	1.68	33.19	31.50	1,20	0.58	1.70	1,20	2.84	32.54	•
415820         W84815         0.63           358046         W94591         0.52           80186         T64192         1.07           454317         AA677165         0.33           768111         AA418988         0.67           299024         N75394         0.46           510136         AA053239         0.01           41300         R5642         0.21           278628         N86203         0.48           20843         H62162         0.21           20848         N74623         0.02           29648         N74623         0.02           359040         W92278         0.83           366795         AA029428         0.48           588159         AA135222         1.78           742671         AA402281         0.66           1031375         AA403769         0.80           225684         AA186335         0.23           28558         N62780         1.35	12.91	12.84	12.78	12.72	12.60	12.58	12.52	12.49	12.49	12.47	12.44	12.37	12.35	12.29	12.20	12.16	12.10	12.04	11.98	11,96	11.96	11.73	11.70	11.63	11.63	11.62	11.47	11.48	11.38	11.35	11.26	11.26	11.23	11.23	
358046 W84815 358046 W94591 80186 T64192 454317 AA677165 768111 AA418988 299024 N75394 510136 AA053239 41300 R56642 278629 N86203 1465883 AA684317 208413 H62162 782279 AA431750 296448 N74623 511233 AA083769 588159 AA13622 742671 AA40281 1031375 AA609138 129862 R17096 813687 AA453769 625684 AA186335 289594 N62780 203488 H55955 742671 AA408745 346815 W74471 1456160 AA862465 433350 AA448672 264449 N21233 786073 AA448672 221828 H92234	8.15	6.66	13.65	4.20	8.45	5.81	0.13	2.68	6.06	8.34	3.96	6.39	0.19	2.21	10.12	5.89	21.58	77.7	7.93	0.12	9.57	10.22	15.85	06:0	3.36	3.53	2.61	73.54	3.38	2.02	0.41	7.40	0.68	4.08	
415820 358046 80186 454317 768111 299024 510136 41300 278629 1465883 208413 782279 782279 782279 782279 782279 782279 742671 1031375 742671 1031375 742671 1031375 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742671 742673 746673 746673 746673 746673	٠		1.07												0.83							0.87	1.35	0.08	0.29	0.30	0.23	6.42	0.30	0.18	9.0	99'0	90:0	0.38	•
				•	_		•							-								•					•	_		_		•		4	
1219 1220 1221 1222 1228 1238 1238 1238 1238 1238	•	•																		129862	813697	625684	289594	203488	785788	346615	1456160	433350	112559	753368	264449	786073	221828	384015	

Table 2-10: Prostate

2580523         N30228         0.87         9,88         11,15         1,11         312.8           627222         AA181438         0.07         0.79         11,09         132         30.94           620462         AA181438         0.07         0.79         11,09         132         30.94           620462         AA18893         1.08         11,08         11,06         12.8         31.46           22051         H7024         0.14         1.56         11,00         1.13         31.46           5062         H7024         0.14         1.56         10.99         0.83         31.48           5072         H7024         0.14         1.53         10.89         0.83         31.48           5072         H702         1.06         1.03         0.83         31.48         31.48           5072         H702         1.06         1.03         0.83         31.48         31.48           5072         H702         1.06         1.03         0.83         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48         31.48	Can	cer Cell (	Lines vs. Norl	mal Prostat 2 of 3 or 10	ic Epithellal ( ) fold or high≰	Cell Line: Ger ar in at least '	mal Prostatic Epithellal Cell Line: Genes upregulated 3 fold 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	e Cancer Cell Lines vs. Normal Prostatic Epithellal Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 cancer cell lines.	her in 3 of 3, c	n 5 fold or high	ner in at least
220602         NOUZE         ANDUS         0.77         (1,09         1.32         30.34         0.0           220602         AAA18498         0.07         11.98         11.08         1.26         31.46         0.0           220602         AAA18498         0.06         7.28         11.05         1.33         31.48         0.0           220602         AA18999         0.06         7.26         11.00         1.13         31.48         0.0           207274         M33550         0.24         0.26         10.39         0.83         31.48         0.0           5082         H1778         0.24         2.62         10.39         0.83         31.48         0.0           28725         M3276         0.24         2.62         10.39         0.83         31.48         0.0           28725         M3276         0.24         2.62         10.39         0.83         31.48         0.0           28726         M3276         0.45         1.69         10.70         0.83         31.48         0.0           28727         M46348         0.42         2.62         10.39         0.83         31.48         0.0           28727         M463				78.0	9.68	11,15	1.1	31.28	1.05	<del>-</del> -	
22,722.         AMBINAS         0.00         11.08         11.06         12.6         31.46         0.0           22,722.         AMBINAS         0.00         11.08         11.05         0.91         31.46         0.0           22,056.8         AMBINAS         0.02         3.51         11.04         1.23         31.46         0.0           2375.8         MARZHA         0.04         1.56         11.00         1.13         31.25         0.0           2877.2         H1733         0.20         3.26         10.99         0.89         30.48         0.0           2897.2         H1733         0.24         1.56         10.09         0.87         30.48         0.0           2897.2         H1734         0.24         1.56         10.09         0.87         30.88         1.01         0.0           2897.2         H1734         0.04         1.34         10.75         1.04         28.68         1.01         0.0	ž	228002	NSUSCO	5 6	0.79	11.09	1.32	30.94	0.99		
626462         AA188999         1.00         11.05         0.91         31.48         0.0           6236462         AA188999         1.00         1.20         3.12         1.10         1.13         31.48         0.0           62374         AA40640         0.56         7.28         1.10         1.13         31.48         0.0           2877.4         RAZ563         0.32         3.26         1.00         0.83         31.48         0.0           50582         H17038         0.30         3.26         1.09         0.83         31.48         0.0           50582         H17038         0.30         3.26         1.09         0.83         31.48         0.0           289725         H17038         0.34         1.06         0.89         0.97         31.48         0.0           289726         H17038         0.89         0.07         0.89         0.09         22.45         0.0           289727         R1389         0.12         1.34         10.75         1.06         0.07         0.09         23.48         0.0           280417         AA463463         0.24         2.53         1.05         0.07         23.48         1.01         0	255	627292	AA181495	5 5	41 98	11.08	1.26	31.46	0.52		-
6225618         AA466649         OSES         7.20         1104         1.23         31.25         0.0           270774         N33550         0.32         3.51         1104         1.33         31.25         0.0           28774         RAZ785         0.30         3.26         10.99         0.83         31.48         0.0           3378         RAZ785         0.24         2.62         10.89         0.80         29.45         0.0           3378         RAZ785         0.24         2.62         10.89         0.05         29.45         0.0           28972         NSB278         0.80         0.26         1.01         0.0	526	626462	AA188999	9 (	2 6	11.05	0.91	31,48	0.76	· 	
270274         N33550         0.32         3.51         1.13         31.52         0.0           28774         R40244         0.14         1.56         11.00         1.13         31.52         0.0           39182         H47038         0.24         2.62         10.59         0.83         31.48         0.0           39187         H5405         0.24         2.62         10.59         0.89         31.48         0.0           28725         H6406         0.66         5.00         10.76         0.67         30.86         1.01         0.0           28725         H84138         0.12         1.34         10.75         1.66         22.36         1.01         0.0         24.6         1.01         0.0         24.6         22.63         1.0         0.0         27.7         30.86         1.0         1.0         0.0         27.7         30.86         1.0         1.0         0.0         27.7         30.86         1.0         1.0         0.0         27.7         30.86         1.0         0.0         27.7         0.0         0.0         27.7         0.0         0.0         27.7         0.0         0.0         0.0         0.0         0.0         0.0	257	823618	AA496949	9.66	07.1	<u> </u>	1.23	31.25	0.63	<b>-</b> -	
28774         RAG244         0.14         1.50         2.45         2.45         2.45         1.51         2.45         2.45         2.45         1.51         2.51	528	270274	N33550	0.32	5.5	5 5	1.13	31,52	0.36		
50582         H17038         0.30         3.26         10.39         0.39         29.45         2.5           31478         R42763         0.24         2.62         10.89         0.80         29.45         2.6           83358         T68445         1.46         15.93         10.89         0.97         30.86         1.01           2262313         H93460         0.46         5.00         10.76         0.61         29.58         1.01           2262313         H93460         0.46         5.00         10.76         0.61         29.58         1.01           226231         H93460         0.46         5.00         10.76         0.62         29.36         1.01           226231         H93460         0.46         5.00         10.76         0.62         29.36         1.01           217261         H93460         0.46         5.00         10.70         1.06         29.36         1.01           217261         H93460         0.24         2.56         10.70         1.06         29.76         1.07           217266         AA2160         0.40         4.27         10.60         1.75         29.77         29.86           252956	259	28774	R40244	 4.	0°.T	8 6	180	31,48	0.67	-	
31478         RAZ763         0.24         2.62         10.59         30.68         101           83358         T68445         1.46         15.93         10.89         30.68         101           289725         N58276         0.83         9.08         10.89         0.97         30.68         101           282531         H93460         0.46         5.00         10.76         0.61         29.36         10.99         30.68         101           280433         H93460         0.46         5.00         10.74         0.92         29.74         10.66         29.74         10.66         29.74         10.66         29.74         10.66         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         10.66         20.75         29.74         20.66         20.73         20.75         29.74         20.66         20.7	280	50582	H17038	0.30	3.26	86.01		29.45	2.58	<b>~</b>	
83358         T68445         1.46         15.93         10.89         50.00         1.07           262313         NISSZT6         0.83         9.06         10.76         0.97         30.86           262313         H99460         0.46         5.00         10.76         1.66         29.36           29251         R41389         0.12         1.34         10.75         1.66         29.36           210413         N47214         0.51         5.45         10.74         0.92         29.74           210413         N47214         0.51         2.56         10.70         1.06         29.74           21042         N47214         0.51         10.68         2.06         0.06         29.74           44387         H06225         0.25         2.63         10.68         2.06         0.06         29.77           74266         AA40482         0.41         4.77         10.60         1.65         29.77         29.78         29.77         30.58         29.77         29.06         29.78         29.78         29.78         29.78         29.78         29.78         29.78         29.78         29.78         29.78         29.78         29.78         29.78         <	261	31478	R42763	0.24	2.62	50.50	3 5	-	0.98	-	
289725         NS9278         0.63         9.08         10.89         0.31         0.00           262313         H99460         0.46         5.00         10.78         0.61         29.68           262313         H99460         0.46         5.00         10.76         0.61         29.58           29251         R41389         0.12         1.34         10.75         1.66         29.36           280413         N47214         0.51         2.65         10.70         1.06         29.70           811785         A4463463         0.24         2.56         10.70         1.06         29.70           44387         H06325         0.25         2.63         10.68         2.06         0.06           742816         AA401482         0.40         4.77         10.60         1.65         29.70           653968         AA219315         0.24         2.53         10.52         0.75         30.56           65396         AA219315         0.24         2.53         10.40         1.13         29.06           65396         AA219315         0.24         2.53         10.42         1.01         29.06           65396         AA219315         0	262	83358	T68445	1.46	15.93	10.89	30.00	10.1 An Or	78.0	-	
252313         H99460         0.46         5.00         10.76         0.81         29.56           29251         R41389         0.12         1.34         10.75         1.66         29.36           29251         R41389         0.12         1.34         10.75         1.66         29.74           280413         N47214         0.51         5.45         10.70         1.06         29.74           281785         AA463463         0.24         2.56         10.70         1.06         29.70           44387         H06525         0.25         2.63         10.68         2.06         0.06         29.70           38517         R51236         0.11         1.13         10.62         0.73         31.04           44387         H06525         0.24         2.53         10.60         1.65         29.17           62968         AA219315         0.24         4.77         10.60         1.65         29.17           62968         AA21931         0.24         4.74         10.45         1.01         29.40           687579         AA21561         0.67         6.34         10.41         1.13         29.40           72633         AA6658 <td>28.2</td> <td>289725</td> <td>N59278</td> <td>0.83</td> <td>9.08</td> <td>10.89</td> <td>A.O</td> <td>9 6</td> <td>- F</td> <td>-</td> <td></td>	28.2	289725	N59278	0.83	9.08	10.89	A.O	9 6	- F	-	
29251 R41389 0.12 1.34 10.75 1.66 29.36 28.36 28.47 10.74 0.922 29.74 280443 N47244 0.51 5.45 10.74 0.922 29.74 280443 N47244 0.51 5.45 10.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 29.70 1.06 1.65 29.17 1.06 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 29.70 1.06 1.06 1.06 1.06 29.70 1.06 1.06 1.06 1.06 1.06 29.70 1.06 1.06 1.06 1.06 1.06 1.06 1.06 1.0	3 6	262343	09760H	0.46	5.00	10.76	0.81	28.68	5. 4.	. 🕶	
292011         NAT214         0.51         5.45         10.74         0.92         29.74           280413         NAT214         0.51         5.45         10.70         1.06         29.70           811786         AA463463         0.24         2.56         10.70         1.06         29.70           811786         AA463463         0.24         2.56         10.70         1.06         29.70           38517         R61236         0.11         1.13         10.62         0.73         31.04           742616         AA401482         0.40         4.27         10.60         1.66         29.17           623968         AA219315         0.24         2.53         10.62         0.75         30.56           623968         AA219315         0.24         2.74         10.40         1.13         28.01           784177         AA436561         0.67         6.34         10.41         1.26         29.40           68779         AA436563         1.77         18.36         10.40         1.13         28.01           730294         AA412512         0.33         3.36         10.26         1.20         29.40           744253         AA466289	\$	20202		0.42	¥.	10.75	1.66	29.36	67.1		
286413         N47214         COL         256         10.70         1.06         29.70         11           811786         AA462463         0.24         2.56         10.70         1.06         2.06         0.06         22.70         11           44387         H05255         0.25         2.63         10.68         2.06         0.06         22.7         0.06         2.06         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         22.7         0.06         0.06         22.7         0.06         0.06         22.7         0.06         0.06         22.7         0.06         0.06         22.7         0.06         0.06         22.7         0.06         0.07         0.06         0.07	265	16767	K41309	; ; ;	5.45	10.74	0.92	29.74	1.56	<b>-</b> •	
811785         AA463463         D.24         2.50         10.68         2.06         0.06         22           44387         H06525         0.25         2.63         10.68         2.06         0.06         22           44387         H06525         0.25         2.63         10.68         2.06         0.06         22           38517         F51236         0.11         1.13         10.62         0.73         31.04         0           742616         AA401482         0.40         4.27         10.60         1.65         29.17         0           623966         AA219315         0.24         2.53         10.52         0.75         30.58         0           687579         AA236561         0.67         6.34         10.41         1.26         29.40         0 <td>266</td> <td>280413</td> <td>N47214</td> <td>5 6</td> <td>99.6</td> <td>07.01</td> <td>1,06</td> <td>29.70</td> <td>1.34</td> <td>-</td> <td></td>	266	280413	N47214	5 6	99.6	07.01	1,06	29.70	1.34	-	
44387         H06525         0.25         4.03         1.04         31.04           38517         R51236         0.11         1.13         10.62         0.73         31.04           742616         AA401482         0.40         4.27         10.60         1.65         28.17           742616         AA401482         0.40         4.27         10.60         1.65         29.17           629968         AA219315         0.24         2.53         10.52         0.75         30.56           587579         AA42658         1.77         18.36         10.41         1.12         28.01           730294         AA412512         0.33         3.36         10.32         1.20         28.15           730294         AA412512         0.33         3.36         10.26         1.12         28.46           745654         AA412512         0.33         3.36         10.26         1.12         28.15           72653         T50397         0.01         0.10         10.26         1.20         28.58           773423         AA460289         0.34         3.44         10.24         1.33         28.32           77551         AA450346         0.42	1267	811785	AA463463	0.24	6.30	40 AB	2.06	0.06	29.91	-	
38517         R51236         0.11         1.13         10.02         0.15         20.17           742616         AAd1482         0.40         4.27         10.60         1.65         29.17           742616         AAd1482         0.40         4.27         10.60         1.65         29.17           55306         AAZ19315         0.24         2.53         10.52         0.75         30.58           687579         AAZ36561         0.67         6.94         10.41         1.26         29.40           730294         AA412512         0.33         3.36         10.32         1.73         28.01           730294         AA412512         0.33         3.36         10.26         1.12         28.46           72653         750397         0.01         0.10         10.26         1.20         28.15           73654         AA406289         0.34         3.44         10.26         1.33         28.36           795723         AA402289         0.34         3.44         10.26         1.33         28.36           773423         AA4028038         0.15         1.53         10.20         1.31         28.36           773423         AA4028038	268	44387	H06525	0.25	7.63	9 6	i c	31.04	60.0	-	
742816         AA401482         0.40         4.27         1050         1.05           623968         AA213315         0.24         2.63         10.52         0.75         30.56           53308         RA5402         0.45         4.74         10.45         1.01         29.08           687579         AA236561         0.67         6.94         10.40         1.13         28.01           784177         AA446658         1.77         18.36         10.40         1.13         28.01           730294         AA412512         0.33         3.36         10.32         1.20         28.15           730294         AA415512         0.33         3.36         10.26         1.12         28.46           1466549         AA485140         0.44         4.50         10.26         1.12         28.16           730294         AA415512         0.01         0.10         10.26         1.20         28.56           755723         AA460289         0.34         3.44         10.24         1.33         28.36           773423         AA460289         0.34         3.34         10.20         1.31         28.36           743516         AA653466         0.42 <td>269</td> <td>38517</td> <td>R51236</td> <td>0.11</td> <td>1.13</td> <td>10.62</td> <td>2 4</td> <td>29 17</td> <td>0.97</td> <td>-</td> <td></td>	269	38517	R51236	0.11	1.13	10.62	2 4	29 17	0.97	-	
629968         AAZ1931\$         Q.24         2.63         10.52         0.75         0.75           35308         RA5402         Q.45         4.74         10.45         1.01         29.08           687579         AAZ36561         Q.67         6.94         10.41         1.26         29.40           687579         AAZ36561         Q.67         6.94         10.41         1.26         28.40           78477         AA44658         1.77         18.36         10.20         1.13         28.01           730294         AA412512         0.33         3.36         10.26         1.12         28.46           1466549         AA885140         0.44         4.50         10.26         1.12         28.46           72663         750397         0.01         0.10         10.26         1.20         28.56           73652         74426038         0.15         1.53         10.20         0.52         28.65           79572         AA426038         0.15         1.53         10.20         1.31         28.36           79570         AA426038         0.15         1.408         10.19         1.38         28.36           79570         AA538945	270	742616	AA401482	0.40	4.27	10.60	3 1	89.00	0.24	•	
35308         R45402         0.45         4.74         10.45         1.01         29.00           687579         AA236561         0.67         6.94         10.41         1.26         29.40           784177         AA446658         1.77         18.36         10.40         1.13         28.01           730294         AA412512         0.33         3.36         10.32         1.20         28.15           730294         AA412512         0.33         3.36         10.26         1.12         28.01           730294         AA412512         0.31         0.44         4.50         10.26         1.12         28.05           7563         750397         0.01         0.10         10.26         1.20         28.36           795723         AA40289         0.34         3.44         10.24         1.33         28.35           795724         AA450389         0.15         1.53         10.20         0.52         29.62           773423         AA450894         0.42         4.33         10.20         1.31         28.36           28000         N51107         1.38         14.08         10.19         1.36         28.36           28108		629968	AA219315	0.24	2.53	10.52	0.75	30.00 80.00	1 26	•	
687579 AA236561 0.67 6.94 10.41 1.26 28.40 784177 AA446658 1.77 16.35 10.40 1.13 28.01 730294 AA412512 0.33 3.36 10.32 1.20 28.15 730294 AA412512 0.33 3.36 10.32 1.20 28.15 7466549 AA865140 0.44 4.50 10.26 1.12 28.46 72663 72663 0.34 3.44 10.24 1.33 28.32 773423 AA460289 0.34 3.44 10.24 1.33 28.32 773423 AA460289 0.15 1.53 10.20 0.52 773423 AA456038 0.15 1.53 10.20 1.31 28.06 795170 AA453466 0.42 4.33 10.20 1.31 28.06 795170 AA453466 0.42 4.33 10.20 1.31 28.06 743519 AA609415 0.53 5.32 10.01 1.20 27.58 898038 AA598943 0.35 5.32 10.01 1.20 27.58 44303 H06380 0.73 7.21 9.97 1.33 28.16 251685 H96738 0.46 6.50 9.84 1.03 28.15 251685 H96738 0.46 4.57 9.87 0.49 28.25 344010 W70242 0.67 6.60 9.84 1.03 27.33		3530B	R45402	0.45	4.74	10.45	1.01	29.00	- 6	•	
784177         AA446658         1.77         18.36         10.40         1.13         28.01           730294         AA412512         0.33         3.36         10.32         1.20         28.15           730294         AA412512         0.33         3.36         10.22         1.20         28.15           1466549         AA485140         0.44         4.50         10.26         1.12         28.46           72653         75473         0.01         0.10         10.26         1.20         28.56           775423         AA460289         0.15         1.53         10.20         0.52         29.52           773423         AA45038         0.15         1.53         10.20         0.52         28.36           795170         AA450486         0.42         4.33         10.20         1.31         28.36           282000         N51107         1.38         14.09         10.30         1.36         28.36           282000         N51107         1.38         14.09         1.36         28.36           283038         AA598943         0.53         5.32         10.00         1.65         28.36           284950         AA65865         0.67	7/7	647570	A A 236561	0.67	6.94	10.41	1.26	29.40	000	• •	
730294         AA412512         0.33         3.36         10.32         1.20         28.15           730294         AA412512         0.33         3.36         10.22         1.20         28.46           1466549         AA885140         0.44         4.50         10.26         1.72         28.46           72653         75037         0.01         0.10         10.26         1.20         28.56           775423         AA460289         0.34         3.44         10.20         0.52         29.52           773423         AA45038         0.15         1.53         10.20         0.52         29.62           795170         AA453466         0.42         4.33         10.20         1.31         28.36           795170         AA453466         0.42         4.33         10.20         1.31         28.36           282000         NS1107         1.38         14.09         1.38         28.36           282000         NS1107         1.38         14.09         1.36         28.36           289038         AA598943         0.53         5.32         10.01         1.65         28.36           289038         AA69636         0.67         6.03	3	200	1000000	1.1	18.36	10.40	1.13	28.01	2.03	- ,	
730294         AAA12512         U.S.S         10.26         1.12         28.46           1466549         AA885140         0.44         4.50         10.26         1.20         28.58           72663         75633         760397         0.01         0.10         10.24         1.33         28.32           795723         AA460289         0.34         3.44         10.24         1.33         28.32           773423         AA45038         0.15         1.53         10.20         0.52         29.62           795170         AA450366         0.42         4.33         10.20         1.31         28.36           282000         N51107         1.38         14.08         10.19         1.38         28.36           282000         N51107         1.38         14.08         10.19         1.38         28.36           282000         N5107         1.38         14.09         1.36         28.36           898038         AA598943         0.35         5.32         10.01         1.65         28.99           898038         AA598943         0.35         5.37         10.00         1.65         28.24           44303         H06380         0.45	1274	784177	AA446638		3.36	10.32	1.20	28.15	1.60	<b>-</b>	
1466549         AA885140         0.44         4.50         10.26         1.20         28.58           72663         750397         0.01         0.10         10.26         1.20         28.58           726723         AA460289         0.34         3.44         10.24         1.33         28.32           773423         AA460289         0.15         1.53         10.20         0.52         29.62           773423         AA450386         0.15         1.53         10.20         1.31         28.06           795170         AA453466         0.42         4.33         10.20         1.31         28.36           282000         NS1107         1.38         14.09         10.19         1.38         28.36           743519         AA608415         0.53         5.32         10.01         1.20         27.58           743519         AA608415         0.53         5.32         10.01         1.65         28.36           898038         AA598943         0.35         3.47         10.00         1.65         28.99           849038         AA658943         0.46         6.03         9.97         1.33         28.24           44303         H06380	1275	730294	AA412512	C.33		40.7B	1.12	28.46	1.20	-	
72663         750397         0.01         0.10         10.24         1.33         28.32           795723         AA460289         0.34         3.44         10.24         1.33         28.32           773423         AA460289         0.15         1.53         10.20         0.52         29.62           773423         AA453466         0.42         4.33         10.20         1.31         28.06           795170         AA453466         0.42         4.33         10.19         1.38         28.36           282000         N51107         1.38         14.09         10.19         1.38         28.36           743519         AA659843         0.53         5.32         10.01         1.65         26.99           88038         AA598943         0.35         3.47         10.00         1.65         26.99           88038         AA628950         0.60         6.03         9.97         1.33         27.67           251685         H96738         0.46         4.57         9.87         0.49         28.25           244010         W70242         0.67         6.60         9.84         1.03         27.33           775540         AA459652	1276	1466549	AA885140	9. 2	00.4	97.0	2.5	28.58	00.	<b>-</b> -	
795723         AA460289         0.34         3.44         10.24         1.53         10.20         0.52         29.62           773423         AA426038         0.15         1.53         10.20         1.31         28.05           795170         AA453466         0.42         4.33         10.20         1.31         28.05           282000         N51107         1.38         14.08         10.19         1.38         28.36           743519         AA609415         0.53         5.32         10.01         1.20         27.58           898038         AA659843         0.35         3.47         10.00         1.65         26.99           898038         AA624459         0.60         6.03         9.97         1.33         27.67           44303         H06380         0.73         7.21         9.92         1.63         28.25           251685         H96738         0.46         4.57         9.84         1.03         28.16           344010         W70242         0.67         4.44         9.84         1.30         27.33           795540         AA459652         0.45         4.44         9.84         1.30         27.33	1277	72663	T50397	0.0	0.10	10.26	5	28.32	1.07	-	
773423         AA426038         0.15         1.53         10.20         0.32           795170         AA453466         0.42         4.33         10.20         1.31         28.06           282000         N51107         1.38         14.09         10.19         1.38         28.36           743519         AA609415         0.53         5.32         10.01         1.20         27.58           898038         AA598943         0.35         3.47         10.00         1.65         26.99           898038         AA629843         0.36         6.03         9.97         1.33         27.67           44303         H06380         0.73         7.21         9.92         1.63         28.24           44303         H06380         0.73         7.21         9.87         0.49         28.25           251685         H96738         0.46         4.57         9.84         1.03         28.16           344010         W70242         0.47         9.84         1.30         27.33           795540         AA459652         0.45         4.44         9.84         1.30         27.33	1278	795723	AA460289	0.3 X	3.44	10.24	3 5	29.62	0.45	-	
795170         AA453466         0.42         4.33         10.20         1.31         28.36           282000         N51107         1.38         14.09         10.19         1.38         28.36           743519         AA609415         0.53         5.32         10.01         1.20         27.58           898038         AA598943         0.35         3.47         10.00         1.65         26.99           898038         AA598943         0.36         6.03         9.87         1.33         27.67           364950         AA024459         0.60         6.03         9.87         1.63         28.24           44303         H06380         0.73         7.21         9.87         0.49         28.25           251685         H96738         0.46         4.57         9.87         0.49         28.25           244010         W70242         0.67         6.60         9.84         1.03         27.33           795540         AA459652         0.45         4.44         9.84         1.30         27.33	1279	773423	AA426038	0.15	1.53	10.20	33. 1.2	28.06	1.21	-	
282000         N51107         1.38         14.08         10.19         1.35         2000           743519         AA609415         0.53         5.32         10.01         1.20         27.58           898038         AA598943         0.35         3.47         10.00         1.65         26.89           898038         AA598943         0.35         3.47         10.00         1.65         26.89           364950         AA024459         0.60         6.03         9.87         1.33         27.67           44303         H06380         0.73         7.21         9.87         0.49         28.25           251685         H96738         0.46         4.57         9.87         0.49         28.25           344010         W70242         0.67         6.60         9.84         1.03         28.16           795540         AA459652         0.45         4.44         9.84         1.30         27.33	1280	795170	AA453466	0.42	4.33	10.20	15.	28.36	0.82	<b>-</b>	
743519         AA609415         0.53         5.32         10.01         1.20         27.50           898038         AA598943         0.35         3.47         10.00         1.65         26.89           364950         AA024459         0.60         6.03         9.97         1.33         27.67           44303         H06380         0.73         7.21         9.97         1.63         28.24           251685         H96738         0.46         4.57         9.87         0.49         28.25           344010         W70242         0.67         6.60         9.84         1.03         28.16           795540         AA459652         0.45         4.44         9.84         1.30         27.33	1284	282000	N51107	1.38	14.09	10.19	1.38	S	1.26	-	
898038 AA598943 0.35 3.47 10.00 1.65 20.39 384950 AA024459 0.60 6.03 9.97 1.33 27.67 44303 H06380 0.73 7.21 9.92 1.63 28.24 251685 H96738 0.46 4.57 9.87 0.49 28.25 344010 W70242 0.67 6.60 9.84 1.03 28.16	1282	743519	AA609415	0.53	5.32	10.01	1.20	90'77	<u> </u>		
384950 AA02459 0.60 6.03 9.97 1.33 27.67 44303 H06380 0.73 7.21 9.92 1.63 28.24 251685 H96738 0.46 4.57 9.87 0.49 28.25 344010 W70242 0.67 6.60 9.84 1.03 28.16 795540 AA459652 0.45 4.44 9.84 1.30 27.33	200	860808	AA598943	0.35	3.47	10.00	1.65	66.97	9	•	
44303 H06380 0,73 7,21 9,92 1,63 28,24 44303 H06380 0,46 4,57 9,87 0,49 28,25 251685 H96738 0,46 6,60 9,84 1,03 28,16 795540 AA459652 0,45 4,44 9,84 1,30 27,33	3	26.4050	AA024459	0.60	6.03	6.97	1.33	19.17		. 🕶	
251685 H96738 0.46 4.57 9.87 0.49 28.25 251685 H96738 0.46 6.60 9.84 1.03 28.16 344010 W70242 0.67 6.60 9.84 1.30 27.33 795540 AA459652 0.45 4.44 9.84 1.30	1 0	44303	HORRAD	0.73	7.21	9.92	1.63	26.24	<b>b</b> 0	• •	
251685 H567.50 5.10 5.60 9.84 1.03 28.16 3.44010 W70242 0.67 6.60 9.84 1.30 27.33 795540 AA459652 0.45 4.44 9.84 1.30 27.33	C97	20074	00001	0.46	4.57	9.87	0.49	28.25	0.60	- ,	
344010 W/V442 0.45 4.44 9.84 1.30 27.33 795540 AA459652 0.45 4.44 9.84 1.30	1286	500167		74.0	999	9.84	1.03	26.16	2.33	- ,	
795540 AA459652 0.43	1287	344010		9	77 7	9.84	1.30	27.33	0.90	-	
	1288	795540		? ? 	ŗ	, נ					

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	-	-	-	-	-	<del>-</del>	-	-	<b>-</b> -		-	-	-	-	-	-	-	-	-	-	-	-	-	-		-	-	-	-	-	-	
0.95	0.81	1.04	57.0	<del>1</del> ,08	26.83	1.31	0.24	1.05	0.43	2.20	1.03	23.15	0.93	0.74	2.61	2.67	0.12	25.05	2.88	1.80	0.49	1.00	0.65	2.23	1.22	1.86	1.78	1.78	1.18	1.23	1.79	0.66	2.72	1.26	
27.38	27.43	26.98	27.37	27.09	0.40	26.66	75.72	26.48	26.99	24.20	26.65	2.81	2.59	. 26.29	23.30	23.87	27.22	1.00	24.01	25,00	25.75	25.66	25.56	24.09	24.82	24.08	2.38	23.59	24.68	24.09	24.14	25.33	23.64	24.08	
1.16	1.15	1.25	1.14	1.02	1.83	1.01	0.74	1.31	1.25	2.24	0.87	2.58	24.67	1.17	2.22	1.48	0.68	1.90	0.96	1.03	1.55	4,00	1.38	1.09	1.18	1.07	22.82	1.61	1.0	1.57	96.0	0.81	0.35	1.26	
9.82	9.80	9.76	9.75	9.73	9.69	99'6	9.65	9.61	9.56	9.55	9.52	9.51	9.40	9.40	9.38	9.34	9.34	9.32	9.28	9.28	9.26	9.22	9.20	9.14	9.07	9.00	8.99	8,99	8.97	96.9	98'8	8.93	8.91	8.87	38
12.27	. 5.41	5.49	7.41	4.86	1.71	1.53	1.56	6.78	4.97	8.52	4.77	33.20	3.37	6.72	0.72	1.69	0.94	0.09	99.2	1.40	0.19	0.09	3.06	2.38	3.63	3.37	19.69	0.91	6.33	0.48	3.83	8.93	0.87	70.0	
1.25	0.55	0.56	0.76	0.50	0.18	0.16	0.16	0.71	0.52	1.00	0.50	3,49	0.36	0.71	0.08	0.18	0,10	0.0	. 0.82	0.15	0.02	0.01	0.33	0.26	0.40	0.37	2.19	0.10	0.71	0.05	0.43	4.00	0.10	0.01	
T97599	W79499	AA873089	AA406213	N52876	H11730	H52446	H05626	H89331	AA432141	AA199717	R61871	AA115076	AA459941	N70654	H23555	R39901	H05741	171042	AA127069	AA705858	W69649	T49325	W74633	AA774645	N35603	R41357	AA426113	AA485429	N27272	AA490058	W70074	AA437107	AA156873	AA035144	
121633	347020	1325751	742874	283619	47793	201931	43422					491565																_		•	343977			_	
1289	1280	1291		_		1295		1297				1301					1306					1311		1313		1315	1316	1317	1318	1319	1320	1321	1322	1323	

static Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prosta

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

785504	4 AA452095	220	17.90	8.15	2.90	0.49	21.07	
•	•	0.36	2.94	8.11	1.08	1.49	21.75	
•		0.24	1.93	8.07	1.00	22.43	0.77	
		0,45	3.62	8.03	1.35	21.23	1.51	
	`	0.57	4.55	8.03	1.14	20.14	2.80	
-		90.0	0.40	8.02	1.92	19.67	2.29	
		0.35	2.83	8.01	0.90	22.05	1.07	
		0.57	4.52	7.98	1.66	21.87	0.42	
		0.03	0.25	7.97	2.14	20.92	0.85	
		0.19	8	2.96	0.99	22.36	0.53	
•		0.65	5,16	7.96	1.05	21.01	1.82	
		25.0	4.12	7.96	20.12	1.68	2.07	
		40	11.10	7.95	1.35	21.75	0.74	
	٩	0.15	1.17	7.92	0.77	22.04	0.96	
Ť		60,	8.58	7.85	1.33	20.70	1.52	
	•	0.74	5,83	7.83	1.09	21.42	0.97	
	•	R <sub>0</sub>	1.78	7.77	0.65	21.62	1.05	
1376 280266		0.53	4.12	7.74	6.73	21.04	<del>.</del> .	
٠	•	0.90	6.90	7.71	1.21	20.67	1.23	
		0.67	5.09	7.63	1.43	20.64	0.82	
		0.01	90.0	7.62	0.56	21.29	1.00	
• •	•	1.49	11.32	7.60	1.33	20.21	1.27	
		0.14	1.07	7.56	1.39	19.58	1.72	
		0.56	4.20	7.56	0.99	20.96	0.71	
•		0.80	6.05	7.55	0.56	21.62	0.48	
	•	90.0	0.46	7.54	2.35	18.03	2.23	
		0.10	0.79	7.53	1.68	18.23	2.69	
•-	•	0.44	3.30	7.49	0.67	1.28	20.54	
		0.05	0.41	7.48	1.30	1.87	19.28	
		9.84	6.24	7.41	1.45	1.27	19.52	
**		0.01	0.10	7.41	0.76	0.44	21.03	
	٩	3.54	26.15	7.36	1.82	0.14	20.18	
	•	0.0	0.07	7.35	2.78	1.00	18.26	
1392 795199	•	0.54	3.92	7.33	990	20.91	0.41	
4	•	97.0	5	7.54	ţ	48.62	200	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

0.19 1.40 0.22 1.32 0.26 0.36 0.36 0.36 0.37 0.37 0.37 0.36 0.37 0.36 0.36 0.36 0.36 0.36 0.36 0.36 0.36
81.02 1.02 1.03 1.03 1.04

ulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prostat

ate Ca	ancer Cell	ate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or l 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	nal Prostation of 3 or 10	c Epithelial C. fold or higher	ell Line: Gen in at least 1	mal Prostatic Epithelial Cell Line: Genes upregulated 3 fold 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines	d 3 fold or high ell lines.	ner in 3 of 3, or	5 fold or 1
4730	489572	AA101840	024	29.	6.78	1.09	16.56	2.69	<b>-</b> .
2 5	58052	T52820	0.0 F	200	6.77	6.	<b>1</b> .8	18.30	<b>.</b>
3 3	593652	AA166703	0,10	0.71	6.77	0.97	18.08	1.24	<b>.</b>
2 5	505227	AA142923	0.38	2,59	6.75	1.20	17.97	4.09	
3 . 5	257167	N30562	0.70	4.72	6.75	0.76	18.29	1.19	<del></del>
3	705820	AA61507	8 9	5.68	6.74	1.07	18.42	0.74	<b>-</b>
3 5	1412398	AA644998	0.38	2.58	6.73	0.92	18.56	0.73	<b>-</b> .
3 5	811048	44485427	0.02	0.11	6.72	0.63	18.92	0.63	<b>-</b>
2 5	1422424	A4827369	0.51	3.42	6.70	0.51	18.93	0.67	<b>-</b>
<u>}</u>	781048	AA46451	77.7	51.77	99.9	1.79	0.83	17.37	-
2 5	376040	AA040332	0.79	5.28	99.9	<b>4.1</b>	16.50	2.35	_
3 5	260216	N32095	1.81	12.09	6.66	0.51	18.64	0.83	<b>.</b>
} }	788234	A A 4 5 4 0 B 0	0.81	5,41	6.65	0.86	18.31	0.76	<b></b> .
į ;	17307	A168278	0.57	3.47	6.63	1.27	18.25	0.38	<del>-</del> -
784	27333	N36270	9.0	1.22	6.62	<del>2</del> 8.	16.69	<u>4.</u>	-
3	2000	NATOTE	72.0	1.82	9.60	1.26	. 15.70	2.84	<del></del>
<u> </u>	1048586	AASOBASS	2.91	19.16	6.59	1.72	0.22	17.84	_
}	ABUTOR	A A S T 7 5 2 2	0.01	0.07	6.58	2.45	<del>1</del> .8	16.28	-
	797978	A4401895	0.35	2.30	6.56	0.65	17.83	1.02	-
Ì	788240	A 4 4 5 4 0 8 2	60.0	0.60	9.56	1.01	18.57	0. <del>1</del>	<b>.</b>
2 3	141061	W58291	0.55	3.63	6.55	1.48	17.20	0.97	-
1	1055144	AA621363	0.69	4.51	6.55	1.90	16.06	1.68	<b>.</b>
2 4	796369	A456148	1.37	8.94	6.54	1.12	18.17	0.34	<b>-</b>
C37	35620	R45832	0.21	1.36	6.53	1,39	17.90	0.30	<del>-</del>
1 2	246897	W78168	0.48	3.15	6.53	0.81	18.01	0.76	-
3 3	1461161	AA868038	0.48	3.12	6.52	16.55	1.51	1.50	<b>-</b>
1455	361656	W96273	0.28	1.81	6.52	2.01	2.64	14.90	<b>.</b>
345	325182	W49619	29.0	4.15	6.51	1.28	17.38	0.87	•-
1447	815051	AA465168	0.86	5.61	6.51	0.84	2.07	16.61	-
347	1417935	A A B 7 B 7 3 9	0.36	6.21	6.50	0.51	18.14	0.84	-
3 2	384939	AA708686	0,28	<b>4</b> 8.	6.48	1.01	17.01	1.43	<b>-</b>
1460	745493	AA625990	0.44	2.83	6.48	16.43	<del>1</del> .	1.17	-
2 2	38528	R51261	0.16	<u> </u>	6.48	1.90	14.61	2.93	-
54	809611	AA458487	0.01	9.04	6.48	0.87	17.11	<del>1</del> .	<b>-</b>
3	290162	N63286	<b>2</b> .	9.86	6.41	£.5	14.87	2.82	_
<u>;</u>					42				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

AAA48270         1.12         7.19         6.39         0.49         0.43           AAA21271         0.30         1.88         6.37         1.80         16.26           W60288         0.57         3.64         6.35         1.27         16.26           AAA63893         0.85         5.43         6.35         0.85         17.18           AAA65704         0.25         1.86         6.29         0.65         16.51           AAA40606         1.15         7.26         6.29         0.65         16.51           AAA41735         0.21         1.33         6.27         1.16         16.50           AAA4885         0.48         2.38         6.28         1.76         16.30           AAA4885         0.41         3.82         6.24         1.16         16.53           AAA4885         0.48         2.38         6.22         0.89         16.75           AAA4885         0.49         6.27         1.16         16.53         16.53           AAA4885         0.41         0.70         6.19         1.76         16.53           AAA4885         0.41         0.70         6.19         1.70         1.44           AAA4	505454	AA156433	1.28	8.15	6.39	1.42	16.90	90 7	•
AA421271         0.30         1.88         6.37         1.60         16.24           WWG288         0.57         3.64         6.35         1.27         16.26           AA43993         0.65         5.43         6.35         0.85         17.16           AA43994         0.65         1.60         6.34         1.25         16.71           AA437040         0.25         1.60         6.31         1.15         16.50           AA47355         0.23         1.45         6.29         0.65         16.71           AA417356         0.21         1.33         6.27         1.16         16.50           AA417356         0.48         2.86         6.22         0.89         16.75           AA417356         0.49         6.21         1.16         16.50         16.50           AA417356         0.41         0.70         6.19         1.87         1.44           AA704435         0.41         0.70         6.19         1.87         1.44           AA704456         0.41         0.70         6.19         1.87         1.44           AA70443         0.41         0.70         6.19         1.87         1.44           AA70	TRORAR	AA448770	1.12	7.19	6.39	0.49	0.43	18.24	- '
WORDS         AAAGS993         0.57         3.64         6.35         1.27         16.26           AAAGS993         0.85         6.35         6.35         1.27         16.26           AAAGS993         0.85         6.34         6.35         0.85         17.18           AAAGS993         0.85         6.29         0.85         17.18           AAAGS993         1.15         7.26         6.29         0.65         16.51           AAAGS993         1.15         7.26         6.29         0.65         16.53           AAA41735         0.21         1.43         6.27         1.18         16.53           AAA41735         0.21         1.33         6.27         1.16         16.53           AAA41735         0.21         1.33         6.27         1.16         16.53           AAA41755         0.21         1.33         6.27         1.16         16.53           AAA41756         0.41         0.70         6.19         1.87         1.44           AAA41776         0.41         0.70         6.19         1.87         1.44           AAA41776         0.41         0.11         1.53         1.14         1.14           <	731040	AA421271	0:30	1.88	6.37	1.80	15.24	2.08	
AA45999         0.85         5,43         6.35         0.85         17.18           AA46706         1.96         12.45         6.34         1.25         16.71           AA49704         0.25         1.60         6.31         1.15         16.50           AA40606         1.15         7.26         6.29         0.65         16.71           AA40735         0.21         1.45         6.28         1.76         16.30           AA41735         0.21         1.33         6.27         1.16         16.63           AA44885         0.48         2.86         6.24         1.16         16.63           AA41885         0.48         2.89         6.27         1.16         16.63           AA41885         0.41         0.70         6.19         1.81         1.70           AA41885         0.41         0.70         6.19         1.81         1.44           AA41866         0.11         0.70         6.19         1.81         1.70           AA41867         0.41         0.70         6.19         1.81         1.44           AA41877         0.41         0.70         6.19         1.81         1.70           AA41877	233	WED288	0.57	3.64	6.35	1.27	16.26	1.53	Ψ,
AAA06705         1.86         12.45         6.34         1.25         16.71           AA40606         0.25         1.60         6.31         1.15         16.50           AA40606         1.15         7.26         6.29         0.65         16.91           R61100         0.23         1.45         6.29         0.65         16.91           AA40866         0.21         1.33         6.27         1.18         16.03           AA40885         0.26         0.24         1.76         16.30           AA40885         0.48         2.86         6.24         1.76         16.33           AA40885         0.48         2.89         6.22         0.89         16.75           AA40866         0.11         0.70         6.19         1.81         1.44           AA70846         0.11         0.70         6.19         1.87         1.44           AA70847         0.41         0.70         6.19         1.87         1.44           AA70847         0.41         0.70         6.19         1.87         1.44           AA70847         0.41         0.70         6.19         1.87         1.44           AA40857         0.41	705754	AA453993	0.85	5.43	6.35	0.85	17.18	1.03	_
AA406068         1.15         16.50           AA406068         1.15         1.65         6.29         0.65         16.91           AA406068         1.15         7.26         6.29         0.65         16.91           AA406068         1.15         7.26         6.29         0.65         16.91           AA41735         0.21         1.33         6.27         1.16         16.03           N81666         0.61         3.82         6.24         1.16         16.53           AA40855         0.48         2.96         6.22         0.89         16.75           AA406860         0.17         1.04         6.21         1.81         1.23           AA704943         0.41         0.70         6.19         1.81         1.44           AA704945         0.41         0.70         6.19         1.81         1.44           AA41744         0.71         0.71         6.	787383	A A 0.46705	1.86	12.45	6.34	1.25	16.71	1.08	_
Addresses         1.15         7.26         6.29         0.65         16.91           R51100         0.23         1.45         6.28         1.76         16.03           Ad41355         0.21         1.33         6.27         1.18         16.03           Ad41355         0.21         1.33         6.27         1.16         16.03           Ad41355         0.24         3.82         6.24         1.16         16.03           Ad40686         0.17         1.04         6.21         1.18         2.34           Ad40688         0.17         1.04         6.21         1.81         2.34           Ad419646         0.11         0.70         6.19         1.87         14.44           Ad419646         0.11         0.70         6.19         1.87         1.44           Ad419646         0.11         0.70         6.19         1.87         1.44           Ad421	973678	A 4 40 70 40	0.25	1.60	6.31	1,15	16.50	1.28	-
AA401355 0.21 1.45 6.28 1.76 15.30 AA41355 0.21 1.33 6.27 1.18 16.03 AA40865 0.46 3.82 6.24 1.16 16.03 AA40865 0.46 3.82 6.24 1.16 16.03 AA40865 0.48 2.98 6.22 0.98 16.75 AA40865 0.41 0.70 6.19 1.87 1.44 AA70943 0.41 2.50 6.17 1.87 1.89 AA41276 0.02 0.15 6.17 1.87 1.89 AA427276 0.03 0.15 6.17 1.17 1.81 AA4754 0.04 2.40 6.11 1.10 1.07 16.28 AA4757 0.40 2.40 6.07 0.69 15.22 AA47579 0.40 2.40 6.07 0.69 15.25 AA47787 0.40 2.40 6.07 0.69 15.22 AA47787 0.40 2.40 6.07 1.59 15.36 AA47787 0.50 0.65 3.94 6.05 1.37 15.22 AA47787 0.50 0.65 3.94 6.05 1.37 15.22 AA47787 0.50 0.65 3.94 6.05 1.37 15.32 HY124 0.52 3.17 6.05 1.32 HY114 0.27 1.59 6.00 2.55 1.81 H10011 0.27 1.59 6.00 2.75 1.81 H10011 0.27 1.59 6.00 2.75 1.81 H10011 0.27 1.59 6.00 2.75 1.81	4000	A A 406069	4.55	7.26	6.29	0.65	16.91	1.31	
NASTRESS         0.21         1.33         6.27         1.18         16.03           NASTRESS         0.21         1.33         6.24         1.16         16.03           NASTRESS         0.48         2.88         6.24         1.16         16.53           AAA16830         0.17         1.04         6.21         1.81         2.34           AAA16830         0.17         1.04         6.21         1.81         2.34           AAA16843         0.11         0.70         6.19         1.87         1.44           AAA18744         0.41         0.70         6.19         1.87         1.44           AAA168735         0.02         0.15         6.17         1.87         1.44           AAA18755         0.18         0.11         6.15         1.25         1.11           HZ545         0.16         6.17         1.14         1.53         1.11           HZ545         0.14         4.98         6.14         1.14         1.53           AA47574         0.34         2.06         6.15         1.25         1.11           HZ580         0.47         0.27         1.14         15.31           AA47574         0.34	2002	2440008	, C	1.45	6.28	1.76	15.30	1.79	_
N91566         0.61         3.82         6.24         1.16         16.63           AA44885         0.48         2.88         6.22         0.98         16.75           AA44885         0.48         2.88         6.22         0.98         16.75           AA016870         0.17         1.04         6.21         1.81         2.34           AA016436         0.11         0.70         6.19         1.87         14.44           AA01643         0.41         0.70         6.19         1.87         14.44           AA01643         0.41         0.70         6.19         1.87         14.44           AA166735         0.02         0.15         6.17         1.87         14.44           AA166735         0.02         0.15         6.17         1.87         1.44           AA166735         0.02         0.15         6.17         1.87         1.44           AA41776         0.34         0.11         6.15         1.53         1.11           HZ3509         1.42         8.74         6.14         1.14         15.91           AA41774         0.34         2.08         6.14         1.14         15.91           AA417372<	47476		2. 5	133	6.27	1.18	16.03	1.60	_
AA448855         O.46         2.86         6.22         0.08         16.75           AA448855         0.46         2.36         6.21         1.81         2.34           AA016980         0.17         1.04         6.21         1.81         2.34           AA016986         0.11         0.70         6.19         1.87         1.44           AA19466         0.11         0.70         6.19         1.87         1.44           AA40473         0.41         2.50         6.17         0.41         0.41           AA42176         0.99         6.12         6.17         0.41         0.41           AA42176         0.99         6.12         6.17         0.41         0.41           AA421776         0.99         6.12         6.15         1.25         16.18           AA43674         0.01         4.98         6.14         1.14         15.91           NB4587         0.81         4.98         6.14         1.14         15.81           AA44774         0.34         2.08         6.11         1.01         15.63           AA44774         0.34         2.08         6.11         1.01         15.63           AA477972 <td>081187</td> <td>•</td> <td>1 5</td> <td>3.87</td> <td>6.24</td> <td>1.16</td> <td>16.63</td> <td>0.92</td> <td>-</td>	081187	•	1 5	3.87	6.24	1.16	16.63	0.92	-
AAA18625         CAUG         6.21         1.81         2.34           AA016980         0.17         1.04         6.21         1.81         2.34           AA016930         0.17         1.04         6.20         0.84         17.00           AA18676         0.11         0.70         6.19         1.87         1.44           AA168736         0.02         0.15         6.17         0.41         0.41           AA421276         0.39         6.12         6.15         1.25         16.18           AA421276         0.39         6.12         6.15         1.25         16.18           AA421276         0.39         6.12         6.15         1.25         16.18           AA421276         0.39         6.13         1.14         15.91           NB4587         0.81         4.38         6.14         1.14         15.91           NB4587         0.87         6.14         1.14         15.91           AA44774         0.34         2.08         6.11         1.01         15.63           AA44774         0.34         2.08         6.11         1.01         15.63           AA47897         0.40         2.40         6.0	20200	COCIAN	. a	2.98	22	0.98	16.75	9.9	•
AA016980         0.77         7.16         6.20         0.84         17.00           AA388757         1.15         7.16         6.20         0.84         17.00           AA194646         0.11         0.70         6.19         1.87         14.44           AA70493         0.41         2.50         6.17         1.87         1.89           AA70493         0.02         0.15         6.17         0.41         0.41           AA421276         0.09         6.12         6.15         1.25         16.16           AA421276         0.09         6.12         6.15         1.25         16.16           AA421276         0.09         6.12         6.14         1.14         16.36           N64597         0.31         4.98         6.14         1.14         15.31           N64597         0.31         6.14         1.14         15.31         16.36           AA47544         0.34         6.14         1.14         15.31         15.31           AA47574         0.34         6.14         1.14         15.31         15.31           AA47582         0.14         0.87         6.16         1.71         15.32           AA4	45198/	AA448855	; ;	3	6.21	1.81	2.34	14,49	-
AA388767         1.13         (.19         0.20         0.20           AA194646         0.11         0.70         6.19         1.87         14.44           AA704943         0.41         2.50         6.17         16.37         1.89           AA704943         0.41         2.50         6.17         16.37         1.89           AA421276         0.99         6.12         6.15         1.25         16.16           AA421276         0.99         6.12         6.15         1.25         16.16           AA421276         0.99         6.12         6.14         1.14         16.18           N64597         0.81         4.98         6.14         1.14         15.91           N64597         0.81         6.14         1.14         15.91           AA435069         1.42         8.74         6.14         1.14         15.91           AA435079         0.90         6.14         1.14         15.91         1.71           AA47544         0.34         2.96         6.13         1.01         1.53           AA475574         0.34         0.87         6.09         1.75         1.73           AA405181         0.48 <th< td=""><td>361587</td><td>AA016980</td><td>) )</td><td>; u</td><td>F 20</td><td>28</td><td>17.00</td><td>0.78</td><td>_</td></th<>	361587	AA016980	) )	; u	F 20	28	17.00	0.78	_
AA194646         0.11         0.70         0.19         1.01         1.01           AA704943         0.41         2.50         6.17         15.37         1.89           AA156736         0.02         0.15         6.17         0.41         0.41           AA421276         0.99         6.12         6.15         1.25         16.18           AA21276         0.99         6.12         6.15         1.25         16.18           AA21276         0.99         6.12         6.15         1.25         16.18           AA21276         0.99         6.12         6.15         1.14         0.41           NB4597         0.81         4.98         6.14         1.14         15.91           AA36063         0.47         2.86         6.13         1.01         15.63           AA447574         0.34         2.08         6.11         1.07         15.63           AA47574         0.34         2.08         6.11         1.07         15.63           AA47574         0.40         2.40         6.09         13.75         1.73           AA405181         0.48         2.91         6.07         1.05         16.28           AA4051	727137	•	<del>1.1</del>	91.7	0.50	. 6	14.44	2.25	_
AA704943         Q41         2.50         6.17         15.37         1.03           AA158736         Q.02         0.15         6.17         0.41         0.41           AA421276         Q.99         6.12         6.15         1.25         16.18           AA421276         Q.99         6.12         6.15         1.25         16.16           HZ7545         Q.18         1.11         6.15         1.25         16.16           NB4597         Q.34         8.74         6.14         1.14         15.91           AA447574         Q.34         2.08         6.11         1.01         1.563           AA447574         Q.34         2.08         6.11         1.07         15.63           AA447574         Q.34         2.08         6.11         1.07         15.63           AA47574         Q.34         2.08         6.11         1.07         16.26           AA47585         Q.40         2.40         6.08         0.71         17.19           AA405181         Q.48         2.91         6.07         1.05         16.27           AA405181         Q.48         2.91         6.07         0.69         15.36           AA4	665033	•	٥ <del>1</del>	0.70	6.19	1.07	d .	1.26	_
AA426735         0.02         0.15         6.17         0.41         0.41           AA421276         0.99         6.12         6.15         1.25         16.18           HZ7545         0.18         1.11         6.15         15.27         1.11           N64597         0.81         4.98         6.14         1.14         15.91           N84597         0.81         4.98         6.14         1.14         15.91           AA480163         0.47         2.86         6.13         1.01         15.83           AA47574         0.34         2.08         6.11         1.01         15.83           AA47574         0.34         2.08         6.11         1.01         15.25           AA47574         0.34         2.08         6.11         1.07         16.26           AA47574         0.34         6.09         1.37         1.71           AA779972         0.40         2.40         6.09         0.71         17.19           AA405781         0.48         2.91         6.07         0.69         16.27           AA405792         0.40         2.40         6.07         0.69         15.36           AA405794         0.0	461420		0.41	2.50	6.17	15.37	80.	201	•
AA421276         0.99         6.12         6.15         1.25         16.18           HZ7545         0.18         1.11         6.15         1.57         1.11           NB4597         0.31         4.98         6.14         1.14         15.91           R35609         1.42         8.74         6.14         1.14         15.91           AA180163         0.47         2.86         6.13         1.01         1.563           AA43673         0.47         2.86         6.11         1.07         15.63           AA447574         0.34         2.08         6.11         1.07         15.63           AA447574         0.34         2.08         6.11         1.07         15.63           AA477873         0.50         3.06         6.11         1.07         16.26           AA477872         0.14         0.87         6.08         0.71         17.19           AA405181         0.48         2.91         6.06         0.71         17.19           AA405181         0.48         2.91         6.07         0.69         15.36           AA405181         0.05         3.94         6.05         0.37         1.37         15.22	591540		0.02	0.15	6.17	0.41	0.41	80.7	- •
H27545 0.18 1.11 6.15 15.27 1.11 N64597 0.81 4.98 6.14 1.14 15.91 R33509 1.42 8.74 6.14 1.63 2.30 AA180163 0.47 2.86 6.13 1.01 1.5.63 AA447574 0.34 2.08 6.11 1.81 15.22 AA447574 0.34 2.08 6.11 1.07 16.26 AA873927 1.82 11.09 6.09 13.75 1.73 AA779972 0.40 2.40 6.07 0.69 16.27 AA405181 0.48 2.91 6.07 1.65 16.43 AA405180 0.65 3.94 6.05 1.37 15.22 AA405150 0.65 3.94 6.05 1.37 15.22 AA427404 0.03 0.16 6.05 0.85 2.37 H10011 0.27 1.59 6.00 2.75 1.81 H10011 0.27 1.59 6.00 2.75 1.81	731050		0.99	6.12	6.15	1.25	16.18	<b>2</b>	
N64597         0.81         4.98         6.14         1.14         15.81           R33609         1.42         8.74         6.14         1.63         2.30           AA47674         0.47         2.86         6.13         1.01         15.63           AA47574         0.34         2.08         6.11         1.81         15.22           AA47574         0.34         2.08         6.11         1.07         16.26           AA47574         0.34         2.06         6.09         13.75         1.73           AA97387         0.182         11.09         6.09         13.75         1.73           AA478972         0.14         0.87         6.08         0.71         17.19           AA405181         0.48         2.91         6.07         0.69         15.35           AA405181         0.48         2.91         6.07         1.65         15.36           AA405181         0.48         2.91         6.05         1.37         15.22           AA405181         0.05         3.94         6.05         1.37         15.22           AA427404         0.03         0.16         6.05         2.52         14.36           H10	162734		0.18	1.1	6.15	15.27	<b>-</b> -	2.06	-
R33609         1,42         8.74         6.14         1.63         2.30           AA180163         0,47         2.86         6.13         1.01         15.63           AA47574         0.34         2.08         6.11         1.81         15.22           AA47574         0.34         2.08         6.11         1.07         16.26           AA47574         0.34         0.50         6.09         13.75         1.73           AA87327         1.82         11.09         6.09         13.75         1.73           AA77397         0.14         0.87         6.08         0.71         17.19           AA405181         0.48         2.91         6.07         0.69         15.36           AA405181         0.48         2.91         6.07         0.69         15.36           AA405181         0.48         2.91         6.07         0.69         15.36           AA405181         0.48         2.91         6.05         1.37         15.22           AA427404         0.03         0.16         6.05         2.52         14.36           H10011         0.27         1.59         1.81         1.81           H10011         0.2	289903		0.81	4.98	6.14	1.14	15.91	<u>අ</u>	-
AA180163         0.47         2.86         6.13         1.01         15.63           AA447574         0.34         2.08         6.11         1.81         15.22           NA5983         0.50         3.06         6.11         1.07         16.26           AA873927         1.82         11.09         6.09         13.75         1.73           AA779972         0.14         0.87         6.08         0.71         17.19           AA405792         0.40         2.40         6.07         0.69         16.27           AA405792         0.25         1.54         6.07         0.69         16.27           AA405792         0.25         1.54         6.07         0.69         16.27           AA405792         0.25         1.54         6.05         1.37         15.22           AA405792         0.65         3.94         6.05         1.37         15.22           AA427404         0.03         0.16         6.05         2.52         14.36           H10011         0.27         1.59         1.67         1.81           H10011         0.27         1.59         1.61         1.81           H10011         0.27	435000		1.42	8.74	6.14	1.63	2.30	14.49	-
AA47574         0.34         2.08         6.11         1.81         15.22           NA5883         0.50         3.06         6.11         1.07         16.26           AA873827         1.82         11.09         6.09         13.75         1.73           AA779872         0.14         0.87         6.08         0.71         17.19           AA405181         0.48         2.40         6.07         0.69         16.27           AA405181         0.48         2.91         6.07         1.59         16.27           AA405792         0.25         1.54         8.07         1.59         16.43           AA405792         0.25         1.54         8.05         1.37         15.22           AA405704         0.03         0.16         6.05         0.35         2.37           H71242         0.52         3.17         6.05         2.52         14.36           H70011         0.27         1.59         6.00         2.75         1.81           H70011         0.27         1.59         6.00         2.75         1.81           H70011         0.27         1.59         1.51         1.51           H70011         0.27 <td>600530</td> <td></td> <td>0.47</td> <td>2.86</td> <td>6.13</td> <td>1.01</td> <td>.15.63</td> <td>1.75</td> <td>-</td>	600530		0.47	2.86	6.13	1.01	.15.63	1.75	-
AA973927         1.82         11.09         6.09         13.75         1.73           AA973927         1.82         11.09         6.09         13.75         1.73           AA405181         0.40         2.40         6.07         0.69         16.27           AA405181         0.48         2.91         6.07         1.59         15.36           AA405181         0.48         2.91         6.07         1.59         15.25           AA405150         0.65         3.94         6.05         1.37         15.22           AA427404         0.03         0.16         6.05         0.85         2.37           H71242         0.52         3.17         6.05         2.52         14.36           H7011         0.27         1.59         0.01         1.92         1.81           H10011         0.27         1.59         6.05         2.52         1.436           H40011         0.27         1.59         6.01         2.75         1.81           H40011         0.27         1.59         6.00         2.75         1.81           H40011         0.27         1.59         1.51         1.51           H40011         0.27	95565		7	2.08	6.11	1.81	15.22	<b>5.</b> 30	_
AA973927         1.82         11.09         6.09         13.75         1.73           AA779927         0.14         0.87         6.08         0.71         17.19           AA779972         0.40         2.40         6.07         0.69         16.27           AA405181         0.48         2.91         6.07         1.59         15.36           AA405181         0.48         2.91         6.07         1.69         16.43           AA405182         0.25         1.54         8.07         1.05         16.43           AA40570         0.05         3.94         8.05         1.37         15.22           AA427404         0.03         0.16         6.05         0.35         2.37           H71242         0.52         3.17         6.05         2.52         14.36           H7011         0.27         1.59         6.01         1.92         1.81           H10011         0.27         1.59         6.00         2.75         1.81           M4585         0.01         0.06         5.98         1.76         15.18	000701		200	3.06	6.11	1.07	16.26	9.0	_
AA778972         0.14         0.87         6.08         0.71         17.19           AA778972         0.40         2.40         6.07         0.69         16.27           AA405181         0.48         2.91         6.07         1.59         15.36           AA405792         0.25         1.54         8.07         1.05         16.43           AA865150         0.65         3.94         6.05         1.37         15.22           AA427404         0.03         0.16         6.05         0.85         2.37           H71242         0.52         3.17         6.05         2.52         14.36           N49181         1.02         6.12         6.01         1.92         1.81           H10011         0.27         1.59         6.00         2.75         1.81           M4585         0.01         0.06         5.98         1.76         15.18	150717		1 83	11.09	6.09	13.75	1.73	2.80	-
AAA0518 0.40 2.40 6.07 0.69 16.27 AA405181 0.48 2.91 6.07 1.59 15.36 AA4056792 0.25 1.54 8.07 1.05 16.43 AA405704 0.03 0.16 6.05 0.85 2.37 AA427404 0.03 3.17 6.05 2.52 14.36 H71242 0.52 3.17 6.05 1.92 13.20 H40011 0.27 1.59 6.00 2.75 1.81 WA47585 0.01 0.06 5.98 1.76 15.18	200320	•	410	0.87	6.08	0.71	17.19	0.35	-
AA405181 0.48 2.91 6.07 1.59 15.36 AA405181 0.48 2.91 6.07 1.59 15.36 AA405792 0.25 1.54 6.05 1.37 15.22 AA427404 0.03 0.16 6.05 0.85 2.37 H71242 0.52 3.17 6.05 2.52 14.36 H7041 0.27 1.59 6.00 2.75 1.81 WA47585 0.01 0.06 5.98 1.76 15.18	007667		0 40	2.40	6.07	0.69	16.27	1.25	-
AA496792 0.25 1.54 8.07 1.05 16.43 AA4865150 0.65 3.94 6.05 1.37 15.22 AA427404 0.03 0.16 6.05 0.85 2.37 H71242 0.52 3.17 6.05 2.52 14.36 H40011 0.27 1.59 6.00 2.75 1.81 WA47585 0.01 0.06 5.98 1.76 15.18	40100		84.0	2.91	6.07	1.59	15.36	1.25	_
AA865150 0.65 3.94 6.05 1.37 15.22 AA427404 0.03 0.16 6.05 0.85 2.37 AA47742 0.52 3.17 6.05 2.52 14.36 H71242 0.52 3.17 6.05 2.52 14.36 H40011 0.27 1.59 6.00 2.75 1.81 WA7585 0.01 0.06 5.98 1.76 15.18	078767	-	36.0	1 54	8.07	1.05	16.43	0.72	_
AA427404 0.03 0.16 6.05 0.85 2.37 AA427404 0.03 0.16 6.05 0.85 2.37 H71242 0.52 3.17 6.05 2.52 14.36 N49181 1.02 6.12 6.01 1.92 13.20 H10011 0.27 1.59 6.00 2.75 1.81 W447585 0.01 0.06 5.98 1.76 15.18	40/20	Ī.	7 6	200	80.8	1.37	15.22	1.56	_
AA427404 0.03 0.19 0.05 2.52 14.36 H71242 0.52 3.17 6.05 2.52 14.36 1.32 14.36 1.92 13.20 13.20 14.0011 0.27 1.59 6.00 2.75 1.81 1.81 1.81 1.81 1.76 15.18	146985	-	6.6	7 7 6	, S	0.85	2.37	14.92	_
H71242 0.52 5.17 0.55 13.20 13.20 13.20 14.0011 0.27 1.59 6.00 2.75 1.81 1.76 15.18	770989	-	3 6	2 Ç	, d	2.52	14.36	1.26	_
N49181 1.02 6.12 6.01 1.94 1.25 H10011 0.27 1.59 6.00 2.75 1.81 W447585 0.01 0.06 5.98 1.76 15.18	214624		0.52	- -	3		13.20	2.90	_
H10011 0.27 1.59 6.00 2.75 7.01 W47585 0.01 0.06 5.98 1.76 15.18	280214	N49181	1.02	6.12	6.01	1.92	07:01	43.43	•
VA47585 0.01 0.06 5.98 1.76 15.18	46715	H10011	0.27	1.59	6.00	2.75	1.61	7:55 60.	•
	324345		0.0	90.0	5.98	1.76	15.18	9. 1.00	-

if Prostatic Epithelial Cell Line; Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Pros

499 838500		AA457528	ķ	2.05	5.98	1.18	15.89	0.86	-
		AA401409	0.79	4.68	5.85	1.32	15.78	0.75	•
	•	NOTATE	0.67	3.98	5.95	0.60	0.01	17.23	~
		0 A 4 7 0 0 2 R	0.38	2.17	5.95	1.28	16.22	0.34 24	~
	•	AA44256	44	2.60	5.93	1.27	15.41	1.12	-
_		AA775405	0.25	1.49	5.93	0.77	14.53	2.48	-
		A448825	0.79	4.69	5.91	1.10	15.90	0.74	~
		T77936	908	12.14	5.91	0.45	0.54	16.74	-
1506 00500		A A R 2 R 1 3 7	0.23	38.	5.90	0.72	15.08	1.90	<del>-</del>
		H24129	0.37	2.18	5.89	17.10	0.16	0.40	
•		A6418004	0.36	2.11	5.88	1.67	13.14	2.85	-
		HOB18	400	0.21	5.88	2.42	14.95	0.28	-
_		0 A 774678	8.44	37.89	5.88	1.55	3.00	13.10	<b></b>
	•	N20704	0.35	2.08	5.87	1.16	15.23	1.22	<del></del>
		A440955	1.18	6.90	5.87	0.79	15.93	0.88	-
	•	MOSAZE	2	6.10	5.87	1.30	15.63	0.67	-
		ACATT24	0.25	1,46	5.86	1.23	15.07	1.27	
		H44707	0.42	2.46	5.81	14.09	2.49	0.87	
•		AA620927	0.37	2.13	5.81	0.88	15.63	0.91	-
=		N48901	0.47	2.71	5.76	1.07	13.52	2.69	-
		N49619	0.28	1.60	5.76	1.36	14.88	1.05	-
		R40780	0.85	4.87	5.75	1.71	99:0	14.81	<b>-</b>
•	-	WARGGG	1.03	5.93	5.74	1.04	15.61	0.56	₩
		N23652	60.	6.22	5.72	1.37	14.85	96.0	-
		AA443058	4.29	24.55	5.72	0.26	16.15	0.77	~
		94177469	90.0	0.32	5.72	1.68	15.29	0.18	<b></b>
	•	Natable	0.13	0.76	5.70	1.4	15.60	0.07	-
		NEOZO	0.55	3.16	5.70	1.10	14.20	1.79	-
		10002	4.80	77.32	5.69	1.08	2.35	13.64	-
	•	1476	109	6.19	5.67	0.88	15.05	<b>5</b> .08	-
, ,	•	HARADS	0.52	2.93	5.66	1.05	14.61	1.31	-
		14461225	0.41	2.33	5.65	1.01	13.08	2.88	<b></b>
	•	118	0.57	3.19	5.63	2.86	11,46	2.58	-
		4 Z	0.0	90:0	5.61	1.00	1.00	14.84	<b>-</b>
• ••	_	203	0.52	2:30	5.59	1.1	14.64	<u>2</u> .	-

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least

																															٠			
•	•	_	•	•	_	_	_	•	•	-	_	•	_	_	-	_	_	-	-	_	_	_	_	-	-	-	-	-	-	-	<u>-</u>	-	-	•
1.39	2.72	1.4	ž	0.43	2.93	0.80	10.87	1.76	9,34	0.54	14.48	0.93	2.71	14.43	79.0	0.98	0.7	1.59	0.69	13.75	10.73 27.00	0.14	0.52	<u>7.</u>	0.22	1.99	0.11	1.81	0.95	1.12	1,40	1.52	12.24	0.60
14.07	12.89	0.45	13.78	14.42	12.33	14.57	2.90	1.81	14.66	15.17	1.00	14.52	12.03	1.14	15.22	14.23	14.78	13.47	14.20	1.33	2.97	13.71	15.03	13.09	14.13	11.67	15.58	1.81	14.19	14.02	13.69	12.04	2.43	14.21
1.31	1.17	14.87	2.63	1.89	1.47	1.33	2.92	13.10	1.63	06:0	1.11	1.1	1.82	0.93	0.59	<u>1</u> 2	0.88	1.35	1.46	1.27	2.64	2.49	0.78	1.86	1.92	2.61	0.59	12.59	<u>2</u> .	1.04	1.07	2.60	1.48	1.30
5.59	5.59	5.59	5.58	5.58	5.58	5.57	5.56	5.56	5.55	5.54	5.53	5.52	5.52	5.50	5.48	5.48	5.48	5.47	5.45	5.45	5.44	5.44	5.44	5.43	5.42	5.42	5.42	5.41	5.39	5.39	5.39	5.39	5.39	5.37
1.02	6.24	<u>4</u>	0.37	2.21	2.46	2.93	7.23	0.42	0.65	5.82	90.0	3.10	0.16	6.39	0.57	3.15	4.04	1.09	4.31	6.45	12.51	0.40	0.22	3.92	0.25	0.33	0.28	0.03	4.66	3.15	3.29	2.17	13.33	1.45
0.18	1.12	0.19	0.07	0.40	0.44	0.53	1.30	90.0	0.12	1.05	0.01	95'0	0.03	1.16	0.10	0.57	0.74	0.20	0.79	1.18	2.30	0.07	9.0	0.72	0.05	90.0	0.05	0.01	98.0	0.58	0,61	0.40	2.48	0.27
AA598975	AA620463	N51002	H06249	AA404356	AA186613	W95414	AA598501	H12338	R28660	N50432	H51549	AA401472	AA401470	AA703147	W73523	AA609284	T96913	H09541	W95409	W92011	H56028	AA447547	AA620472	N90514	AA009763	AA459702	R36989	AA460328	N48078	N47425	AA176607	AA205803	AA496544	H89713
898288	951102	281243	44287	758370	625616	358212	898218	148469	133864	280633	179753	742589	742590	434845	344505	1031662	121409	45922	358200	415264	203544	782611	951101	306121	429773	795585	27098	795788	281756	280672	609052	645332	755904	240273
1534	1535	1536	1537	1538	1539	1540	1541	1542	1543	154	1545	1546	1547	1548	1549	1550	1551	1552	1553	1554	1555	1558	1557	1558	1559	1560	1561	1562	1563	1564	1565	1566	1567	1568

regulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prost

tate Ca	ncer Cell	Lines vs. Non	mal Prostal 2 of 3 or 10	tic Epithelial of fold or high	mal Prostatic Epithelial Cell Line; Genes upregulated 5 residents of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines	nes upregura 1 of 3 cancer	cell lines.		istate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upreguiace Cancer Cell Lines. 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.
į	60776	7,50,024	5	0.05	5.36	2.88	10.35	2.86	<b>-</b>
6961	86239	AA304611		2.10	5.36	1.17	13.41	1.50	<b>.</b>
0/61	767/71	A 600005	, 50 . 70 . 70	1.33	5.34	13.49	1,23	1.31	<del>-</del>
1571	904705	CEORROAN	2 2	8	ž,	0.99	13.90	1.13	<b>-</b>
1572	562158	AA211565	C./.0	3 6	34	1.08	13.96	0.97	<b>-</b>
1573	270766	N29796	0.0	20.0	5.32	1.04	14.31	0.61	-
1574	742887	AA406220	7	8 86	5.32	0.65	0.18	15.12	-
1575	128058	R09729	9. G	37.	5.31	0.68	14.89	0.38	<b>-</b>
1578	346359	W74216	08.0	9 9	A 22	0.62	13.87	1.43	<b>-</b>
1577	489535	AA098867	1.87	78.8		0.75	14.20	0.97	-
1578	297878	N70038	3.78	20.08	2 6	9 9	14.65	0.76	-
1579	84044 44044	AA485714	0.82	¥.74	ر د د	000	33 +	13.96	-
1580	251591	H86643	0.03	0.18	5.31	0.30	8 5	200	•
1581	745072	AA626275	0.35	1.87	5.30	12.38	04.	5 6	٠ -
1582	378242	AA777024	2.14	11.30	5.29	1.37	13.86	8 6	٠.
4 6 6 6	42120	859618	0.39	2.08	5.29	0.85	14.51	n (	- ,
2 3	278188	MEGORS	1.09	5.76	5.28	2.04	11.84		<b>-</b> ,
2	200017	4 4 4 4 4 8		2.96	5.28	1.40	13.61	0.83	<b>,</b>
1585	000/20	William Milliam	2 6	3.67	5.28	14.20	1.16	0.47	-
1586	124148	R01739	2 5	5 6	5.28	14.23	0.22	1.38	-
1587	757222	AA496149	0.0	, u	4.27	1.12	13.86	0.83	-
1588	595623	AA167338	<b>6</b> .	27.0	, u	, Ç	13.60	1.05	
1589	743452	AA609368	<b>5</b> .0	4.41	0.40	) · ·	53.45	8.	-
1590	454698	AA677200	0.56	2.92	5.25	07.1	2 7	14.83	_
1591	280412	N47208	0.02	0.13	5.24	0.49		-	
1502	360644	AA015819	1.79	9.37	5.24	0.59	13.78	t 6	- •
1503	758309	AA404269	1.08	29.9	5.23	0.33	12.98	R. 7	• •
1504	462963	AA682439	0.43	2.23	5.23	1.88	12.77	3 7	- ,
102	565154	AA195668	44.0	2.30	5.23	1.87	12.97	<b>9</b> 80	- 1
7 6	500703	A A 464 466	220	1.16	5.22	0.13	15.02	0.51	<u>.</u>
20 1	184760	AA161400	, C	76.0	5.21	0.41	12.55	2.66	-
/RGL	0.00	***********	2 6	2.50	5.20	1.26	11.43	2.91	••
886	671706	100603	; ;	0.56	5.20	0.63	14.87	60.0	-
1988	/CI #07	NZUGUZ		2.	5.19	15.	12.74	1.28	<b>-</b>
9	283890	N20/8/	20.0	<b>?</b>	7.18	14.21	0.93	0.40	<b></b>
1601	51011	H19242	2	30.0	. 4	4.	12.97	1.40	-
1602	162533	H27986	2.59	13.38	5 5	5 E	12.99	1.80	-
<del>2</del>	376435	AA039713	<del>-</del> :	5.75	5.17	2	3		
	,				<b>&amp;</b>				

Table 2-10: Prost

					•	•	1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1. 1	thor in 3 of 3	or 5 fold or higher in at least	بد
ostate Ca	ancer Cell	Lines vs. No	rmal Prostal 2 of 3 or 10	tic Epithelial	Cell Line: Ge er in at least	mal Prostatic Epithellal Cell Line: Genes upregulated 3 fold 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.	ed 3 fold of life cell lines.		ostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3 of 3 of 10 fold or higher in at least 1 of 3 cancer cell lines.	
;	,		76.0	1.00 138	5.17	1.17	12.54	1.80	-	
1604	14/38/2	AA8/29/9	7.0	12.87	5.17	0.92	13.61	96.0	<b>-</b>	
1605	742898	AA405809	2 F	500	5.16	1.37	12.96	1.16	-	
1608	1031736	AA609594	0.73	0.00	, u	1.80	13.19	0.50	-	
1607	53081	R16259	0.69	2 2 3 4 5 6		0.43	14.23	0.82	-	
1608	1470278	AA866029	0.51	2.53	9. i	1 12	13.32	1.02	-	
1609	767414	AA417915	0.59	3.04	ų, i	1.15	12.49	128	-	
1610	341065	W58191	0.52	2.68	5,15	, e. e	7.33	2.19	-	
1611	233712	H79035	0.42	2.14	5.14	26.01	8.5	107	-	
1612		AA428659	0.63	3.22	5. 4	1.32	13.03	8		
1613	•	AA608974	0.71	3.64	5.13	0.80	13.91	8 6		
1614		R92281	1.98	10.18	5.13	10.73	2.45	1 27	- <del>-</del>	
40,4		NA0976	==	5.71	5.13	1.08	13.20	01.1		
1010	•	AA707815	0.27	1.36	5.13	10.73	1.75	2.91		
0101		A 4 0 4 7 7 7	2.39	12.23	5.12	11.35	2.00	2.00	<b>,</b> ,	
761		201015	0.43	2.19	5.11	1.25	13.11	0.98	<b>-</b>	
8191		N46590	90 +	15.	5.10	0.66	13.43	1.23	<b></b>	
1619		44466064	5 6	0.87	5.10	1.51	13.26	0.52	•-	
1620		N52030		424	5.10	1.21	13.06	1.01	<b>~</b>	
1621		AA400389	3 5	a t	70.5	1.55	12.47	1.21	<b>~</b>	
1622		N49109	. 4. . 4.	6. 6	. v	1.14	13.28	0.75	_	
1623	-	N91175	2.45	6.21	, u	2.38	11.90	0.88	. <del></del>	
1624		AA460963	0.35	07:	, u	0.60	13.49	1.08	-	
1625		N45223	0.85	4.51	g w	9.60	10.93	1.62	-	
1626	_	AA455096	0.45	2.25	5.03	20.7	11 65	1.78	-	
1627	278572	N66178	2.84	14.29	0 1	) - C	11.47	1,09	-	
1628	257746	N30597	0.82	4. 1.	20.6	7.00	12.41	0.93	-	
1629	257779	N26993	0.52	2.63		 	00	1.00	•	
1630	362795	AA018683	5	0.03	0.0	2.5	14 04	66	<b>-</b>	
1631	290680	N67619	0.55	2.76	5.00	1.2.1	11.31	56.0	-	
1632	119289	T98056	070	0.98	5.00	1.5	15.15 13.08	1.15	•	
1633	3 285680	N67570	1.14	5.69	4.99	S : 0	5	1.33	-	
<b>23</b>	784130	AA432075	1.61	7.99	4.97	4.51	5 ¢	2	-	
1635	345118	W72920	0.45	2.23	4.96	0.78	( <del>1</del> . 4)	86.0		
1636	428298	AA004946	0.76	3.75	4.95	93. ·	12.00	) (		
1637	913242	AA455804	77	1.17	4.94	0.43	13.07		•	
1838			5.50	27.18	4.94	0.98	2.40	2	-	
					47					

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	R67259	0.31	1.54	4.94	0.53	13.93	0.36	- •
AA156112		0.68	3.33	4. 4.	0.78	12.51	ZC: 1	<b>~ ,</b>
W79445		2.28	11.24	4.93	0.21	14.40	0.20	-
AA406083		0.50	2.47	4.93	1.19	12.16	<del>1</del> .	~
AA454652		2.37	11.69	4.93	1.24	11.25	2.29	-
AA486864		1.30	6.40	4.92	1,30	12.51	0.36	-
1466633 AABB3680	٠	0.16	0.81	4.92	1.09	10.94	2.74	-
		1.12	5.51	4.92	2.39	10.12	2.25	-
N35489		0.19	0.95	4.91	0.82	13.48	0.44	-
AA167500	_	0.68	3.35	4.91	1.14	12.92	0.66	-
H11071		0.23	1.12	4.91	1.98	12.52	0.22	-
AA412247		96.0	4.63	4.90	0.99	13.24	0.48	~
AA101632		0.24	1.16	4.90	14.47	0.18	0.05	-
H15695		0.24	1.19	4.89	1.09	1.08	12.48	-
N50563		50.0	0.19	4.88	1.56	12.12	96.0	-
AA421054		0.27	1.31	4.88	1.60	11.90	<del>1.</del>	<b>-</b>
AA010388		1.14	5.56	4.88	1.59	12.60	0.45	-
AA418744		1.40	6.82	4.87	2.38	0.46	11.76	-
AA146826		0.37	1.82	4.86	1.34	11.70	1.55	<del></del>
AA044903		<b>28</b> .	8.97	4.86	0.79	12.99	0.81	•
W37753		2.68	13.01	4.86	1.09	1.94	1.55	<del></del>
H70775		0.13	0.65	4.86	1.07	11.25	2.25	-
R88267		0.01	0.05	4.86	1.00	1.00	12.57	-
R62470		0.43	2.06	4.85	1.10	12.19	1.25	-
T86932		79.0	3.08	4.84	1.16	12.34	<del>7.</del> 2.	-
R89700		1.7	8.56	4.84	1.76	1.07	11.69	-
N64503		9.64	3.08	4.84	1.04	11.69	1.78	-
AA417275		90.0	0.36	4.83	1.64	12.71	0.13	-
H93328		0.42	2.05	4.83	0.83	1.93	11.72	-
W85709		0.67	3.23	4.82	1.36	11.63	1.48	-
AA045574	_	2.68	12.92	4.82	1.96	11.76	0.74	-
AA489581		0.49	2.35	4.82	76.0	11.67	1.92	-
AA609334		1.12	5.41	4.82	0.99	12.76	0.71	-
97777N		0.0	0.21	4.82	1.07	11.91	1.48	-
AA780190	_	3.28	15.82	4.82	10.78	1.98	1,68	-
				84				

a 3 fold or higher in 3 of 3, or 5 fold or higher in at least Table 2-10: Prosta

;	706100	A A 464110	0.82	3.98	4.81	0.92	12.21	1,32	<del>-</del>
9/4	130133	Dareer Dareer		9	4.79	11.66	1.36	1.36	-
0		0.000	, Y	25.60	4.79	1.89	1.32	11.14	-
9/9	3	8/6/20	} •	8	4.78	2.18	0.33	11.85	-
122	898195	AA598567	5. E	0.30	478	1.67	1.75	10.93	-
878	295483	W05026	54.50	790.007	o ř	- T	12.91	0.38	-
679	429202	AA005358	0.42	2.01	? ;		200	0.29	-
089	703581	AA278759	4.41	21.07	77.4	2 !	200	4 79	-
1681	377051	AA057620	0.14	0.65	4.77	0.47	14.09	2 7	•
683	1390880	AA84447	1.17	5.60	4.77	0.94	12.67	, ;	- •
	743536	A A 609422	0.07	0.31	4.75	2.16	10,53	1.56	-
3 3	705360	A A 450057	0.77	3.40	4.75	0.91	12.47	0.86	-
5	2008/	. 6730024		3.45	4.74	0.47	13.16	0.60	<del></del>
683	0,000	AA118040		5 F F	4.74	1.77	11.33	1.12	-
989	7/8/7	20004	76.0	3	, F	07.0	11.93	1.55	-
1687	289794	N59295	1.45	0.85	2 6	9 9 0	12.95	0.57	-
1688	297857	N68928	0.28	<u> </u>	7.7	7.3	44.28	1.27	-
689	743901	AA634482	0.43	2.04	<b>4</b> .7	ñ i		0.60	•
1690	788519	AA452578	1.20	5.63	4.71	57.	67.21		. •
1691	811907	AA454654	0.10	0.47	4.71	1.82	11.86	7 8	- •
692	796170	AA461086	0.83	3.89	4.70	0.87	12.35	9 4	- •
603	782208	AA431975	990	3.20	4.69	1.26	11.65	<u> </u>	- ,
769	256907	9600EN	0.58	2.71	4.68	12.24	0.73	1.07	- •
808	327480	W20462	0.20	0.92	4.68	1.83	11.61	0.50	
2 4	306513	Notali	1.15	5.40	4.68	0.40	. <del>.</del>	12.09	_
	CERTAR	A 4 10 E E 5 4	=	5.18	4.67	0.54	12.86	0.62	-
5	36067	- COSTA	190	3.00	4.67	1.66	11.17	1.17	-
0 0	4330CA	0.0011507	1.17	5,43	4.65	0.88	12.67	0.41	-
860	10007	A44406		75	4.65	2.12	0.35	11.48	-
3 3	0.1261	AA417833	; q	0.82	4.61	0.90	11.08	<del>2</del> .	-
5	1627	189004	2 6	, r	461	1.05	11.59	1,18	-
702	27/063	DRCREN	. c	; ;	4.60	1.39	11.34	1.08	-
93	26.25	A4401436	S 4	0 4	4.60	1.72	1.15	10.93	-
\$ }	100001	_	2 2	75.6	4.60	1.65	11.22	0.94	-
3	176760	•	73 67	103.69	4.59	1.67	0.01	12.11	-
8	CO00/*!	_	) e	4.74	.58	1.32	11.01	1.41	-
è	25.55	C0471 0404			8.5	1.45	11.30	1.00	-
42			7	5	>>.				

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	_		-	-	-	-	-	~	-	-	-	-	-	<b>-</b> -	-	Ψ-	-	~	-	-	-	-	-	-	-	-	-	<b>—</b>	•-	-	-	-	
1.70	1.05	1.26	1.25	0.28	0.45	<u>추</u>	11.60	2.11	2.26	2.26	0.51	0.44	0.98	1.64	1.50	1.29	0.76	11.48	0.39	1.39	0.71	4.00	2.45	0.87	0.63	2.19	2.	0.77	1.31	1.03	0.21	1.12	0.90	0.48	
11.01	11.06	11.38	11.52	12.54	12.10	11.03	0.90	0.97	10.76	10.02	12.14	12.07	11.61	10.60	11.03	10.50	11.58	1.00	12.03	10.20	12.07	0.61	10.04	11.55	11.78	10.29	10.92	11.30	10.60	11.12	12.55	11.25	11.45	0.39	
0.97	1.53	96.0	0.84	0.79	1.05	52,	1.04	10.45	0.52	1.26	0.88	1.01	0:30	1,24	0.93	1.68	1.12	1.00	1.04	1.80	0.58	11.75	0.86	0.91	0.91	0.82	1.34	1.21	<u>.</u> Æ	1.10	0.49	0.88	0.89	12.36	
4.56	4.55	4.54	4.54	4.54	4.53	4.53	4.51	4.51	4.51	4.51	4.51	4.51	4.50	4,49	4.49	4.49	4.49	4.49	4.48	4.46	4.45	4,45	4.45	4.44	4.44	4.43	4.43	4.43	4.42	4.42	4.42	4.41	4.41	4.41	20
3.82	2.28	2.94	3.70	4.00	4.48	4.06	4.46	1.66	2.99	1.40	3.80	2.04	09'0	2.03	2.17	3.23	4.65	0.04	6.20	8.04	2.59	2.85	2,32	3.08	1.74	0.64	1.90	1.88	1.11	4.58	4.12	1.25	5.27	6.26	
9	0.50	0.65	0.82	0.88	0.99	0.90	66.0	0.37	99'0	0.31	0.84	0.45	0,13	0.45	0.48	0.72	1.01	0.0	1.38	1.80	0.58	0.64	0.52	09.0	0.39	0.15	0.43	0.42	0.25	<b>2</b> .	0,93	0.28	1.19	1.42	
N90583	AA609430	H97614	AA454844	W92798	AA702788	AA487046	AA703117	H17022	AA486288	AA634479	AA454085	AA398247	R46700	W79525.	AA666087	AA101833	N22210	N63768	W86832	AA464861	AA682549	AA777098	R98407	N51574	H11631	T58430	AA453261	N59870	AA085759	AA412499	AA427947	R39014	R22024	N31952	
306243	743531	251212	809976	418400	448024	841153	434828	50781	842847	743893	788246	726663	36491	347066	859382	489549	254468	293013	416889	789383	431255	377898	200656	280431	48060	75644	785349	284803	488276	730147	773512	24895	130742	259374	•
1709	1710	1711	1712	1713	1714	1715	1716	1747	1718	1719	1720	121	172	1723	1724	1725	1726	1727	1728	1729	1730	121	1732	1733	\$	1735	1736	1737	1738	1739	1740	1741	1742	1743	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

	, '	4.41	2.15	0.21	10.87	•	
0.21			) 34 76 O	1.71	10.55		
5.48	-	04.4	10.0	0.18	12.75	<b>-</b>	
•	•	٠. دې	, 5, 6	0.11	10.97	•-	
0.09		4.30	2.12	0.08	10.93	-	
3.85 . 15.86	•	4.30	86.0	11.12	1.00	-	
0.16 0.68	•	5.4 Fe 2	85.0	10.89	1.23	-	
0.19 0.83	•	7. Y	10.31	1.12	1.63	-	
0.47 2.03		) i	1 46	10.22	1.37	-	
4.05		4.35	0 0 0 0 0	10.88	0.78	-	
		δ		11.31	0.62	-	
2.37		ος. 4 Ο		10.36	1.5	-	
1.03	•	4. 34	21.1	9 9	22	_	
0.81 3.50	•	4.34	55.	10.40	3.	•	•
0.17 0.73		4.33	0.81	11.8/		. •	
0.49 2.12		4.33	1.18	70.71			
0.66 2.84		4.32	0.63	EG: 1	6.63	- +	
0.29 1.27	Ĭ	4.32	0.99	10.57	6.5	- <b>•</b>	
0.15 0.65	•	4.29	-03	10.98	0.92	- •-	
0.56 2.39	•	4.29	<u>.</u>		6.9		
0.55 2.36		4.28	0.87	0.30	1.02	-	
1.25 5.35		427	<u>ج</u>	\$ 5	69.0	-	
4.38		4.27	2.01	3 5	7870	-	
4,69		4.26	4 4	10.31	0.68	-	
6.00		4.26	6 6	? = =	0.76	-	
		62.4	1 83	10.13	0.79	<del>-</del>	
5.56		3	3 5	10.76	0.96	-	
N34892 0.65 2.74 4.24		4.24	707	10.79	0.30	_	
		4.74	<u> </u>		11.65	-	
1.10 4.66		5	0.41	5 6	11 10		
11.90		4.21	0.82	0.72	2.5		
0.20 0.86		4.21	1.17	10.21	9 8	- •	
1.98		4.21	1.25	10.85	76'0		
0.77		4.21	1.14	10.41	1.07	_	
07.0		4.20	11.47	0.38	0.75	-	
2.71		4.20	1.18	10.46	96.0	<b>-</b> -	
		51					

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 2 of 3 or 10 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	<b>-</b>	Ψ-	-	-	-	<del>-</del>	-	-	-	-	-	-	-	-	<b>-</b>	-	-	-	-	-	-	-	-	-	-	-	-		<b>-</b>	-	-	<b>-</b>	
0.52	1.12	1.07	0.49	0.81	12.16	0.91	0.89	0.88	0.97	0.80	10.41	1.75	1.05	1.35	0.82	0.76	0:39	0.31	10.18	0.53	10.22	0.23	1.26	11.28	0.28	92.0	0.91	1.03	1.25	0.72	0.56	0.55	0.91	0.62	
10.99	0.35	10.39	0.51	0.02	0.15	10.52	10.31	11.10	10.02	10.33	1.00	10.14	10.78		10.57	10.58	10.32	10.47	1.18	10.08	0.02	11.56	10.38	0.27	11.02	10.39	10.21	10.27	10.00	10.15	10.51	10.42	10.16	10.43	
1.08	11.10	1.10	11.57	11.74	0.25	1.12	1.35	0.54	1.41	1.25	0.93	0.41	0.47	0.89	0.89	0.93	0.95	1.43	0.85	1.60	1.93	0.28	0.52	0.59	0.82	0.86	0.96	0.75	0.79	1.16	0.93	96.0	0.80	0.80	
4.20	4.19	4.19	4.19	4.19	4.19	4.18	4.18	4.17	4.13	4.13	4.11	4.10	4.10	4.10	4.09	4.09	4.09	4.07	4.07	4.07	4.06	4.06	4.06	4.04	4.04	4.04	4.02	4.02	4.01	4.01	<b>6</b> .00	3.98	3.96	3.85	52
3.49	5.55	1.75	6.49	2.67	38.53	3.65	4.43	0.35	3.11	4.41	0. 20.	1.03	2.80	2.10	2.74	2.71	1.43	0.13	2.12	6.21	8.26	3.16	14.17	3.89	3.07	6.18	1.71	1.08	0:00	2:00	8.94	1.14	2.21	6.19	
0.83	1.33	0.42	1.55	0.64	9.21	0.87	1.06	0.08	0.75	1.07	0.0	0.25	0.68	0.51	0.67	99.0	0.36	0.03	0.52	1.53	2.04	0.78	3.49	0.96	0.76	1.53	0.44	0.27	0.22	0.50	2.24	0.29	0.56	1.57	
AA453465	N91003	AA412446	NA7941	W95082	AA620995	AA421001	W81117	AA449686	AA455130	AA470079	T50995	AA778533	W80791	AA455358	AA707753	R05660	AA425749	AA479912	H20046	AA447561	AA487146	AA292054	AA187287	AA701933	R15891	AA443823	AA399633	H08560	AA621478	AA136071	N78903	AA194189	R37780	AA159613	
795168	306066	730100	250217	415145	1056172	731298	415413	785760	809863	730406	76647	1048872	347268	812222	412983	29397	773392	772916	172785			725365		435573				45542	1055278	490071	300024	665738	27072	593218	
1779	1780	1781	1782	1783	1784	1785	1788	1787	1788	1789	1790	1791	1792	1793	1794	1795	1796	1797	1798	1799	1800	1801	1802	1803	1804	1805	1806	1807	1808	1809	1810	1811	1812	1813	

Table 2-10: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 3 fold or higher in 3 of 3, or 5 fold or higher in at least 1 of 3 cancer cell lines.

-	-	-	-	-	-	-	-	-	<b>~</b>	<b>-</b>	-	-	-	-	-	-
<b>4</b> .8	0.79	0.22	0.68	10.05	0.14	0.43	0.67	10.13	10.44	0.74	0.48	9.64	0.40	0.02	0.25	0.32
10.07	. 67.0	11.14	10.25	0.95	10.29	10.27	10,18	0.61	0.20	10.26	10.60	10.08	0.12	0.80	10.04	10.14
0.76	10.25	0.46	0.87	0.74	1.30	0.88	0.68	0.78	0.79	0.42	0.25	0.56	10.68	10.29	0.65	0.32
3.94	3.84	3.94	3.94	3.91	3.91	3.86	3.84	3.84	3.81	3.81	3.78	3.76	3.72	3.70	3.65	3.60
2.48	0.05	80	3.36	86	0.77	10.36	0.92	0.74	0.19	0.93	0.42	4.06	24.47	1.73	0.15	0.11
633	200	25.	. 89.0	2.55	0.20	2.69	0.24	0.19	0.05	0.24	0.41	1.08	6.57	0.47	70	0.03
A A ED 404	A 4 4 000 F	00001474	AACOORE	AA470143	HURSOR	AA863169	R60717	AAAGUZBE	AA777886	790967	D80705	AA885470	AARJERGD	AA476867	AACITER 7	AA148735
704607	2000	108711	104103	764111	45783	1455671	42115	705738	440438	757238	1107	1466416	744044	E00555	ACAGCA:	504761
7,07	4 6	619	2 5	2 9	0 0	\$ C82	183	40.4	3 5	3 5	470	1976	1020	190	070	1830

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines

Order	IMAGE Clone ID	Gen Bank Accession Number	MPM1(PrEC)	Ave Cancer Cell Lines Values	Fold(AveCell line/MPM1)	Fold(MPMZ/MP   M1)	Fold(MPM2/MP Fold(MPM272M # of Cancer M1) PM1) Cell Lines	Fold(MPM272/M PM1)	# of Cancer Cell Lines .
_	757165	AA443950	0.0	5.71	591.67	7.33	66.37	1701.31	89
7	592707	AA160608	0.01	1,66	234.30	8.10	680.66	14.13	က
n	220851	H95633	0.01	1.28	196.56	8.10	570.75	10.82	<b>6</b> 2
*	782804	AA448186	0.01	1.81	180.70	9.88	526.08	7.15	ო
10	1343732	AA725564	10.0	1.12	112.48	13.45	301.13	22.85	က
9	784296	AA447079	0.01	0.66	. 02:30	45.84	106.60	43.45	က
7	796569	AA460463	0.01	0.35	63.57	16.10	138.91	35.70	m
æ	1558655	AA976561	90.0	4.36	54.32	97.06	17.92	47.97	က
0	743016	AA406036	0.01	0.50	50.47	15.70	123.80	11.93	6
5	39442	R51617	0.10	5.02	49.93	99.52	44.71	5.49	60
=	33076	R44048	0.80	36.21	45.35	56.36	29.19	50.51	m
12	626390	AA189106	0.07	3.16	45.16	91.62	20.97	22.90	m
13	128083	R09747	0.01	0. 4	44.34	25.03	101.07	6.93	ო
4	812170	AA456035	0.01	0.30	37.18	16.25	60.79	14.51	•
15	50288	H17888	0.01	0.32	36.79	12.25	87.45	10.68	60
16	243181	H94482	0.01	96.0	35.62	5.91	88.28	12.68	•
. 4	256911	N3009B	0.01	0.25	34.44	20.19	73.72	9.41	•
48	1030613	AA608832	0.01	0.33	33.18	5.81	82.72	11.01	гэ
19	487861	AA045436	0.02	0.59	32.57	29.44	39.44	28.82	60
23	270626	N33331	0.01	0.32	31.52	8.48	9.21	76.87	•
23	291057	N72115	0.23	6.85	29.19	24.48	38.30	24.80	60
23	1455835	AA863282	0.01	0.40	29.16	9.10	30.24	48.14	m
23	108378	177729	0.01	0.41	29.13	7.73	36.47	- 43.19	ဗ
24	1391682	AA789328	0.02	2.0	28.83	6.99	65.78	14.72	က
83	263014	H99813	0.63	17.34	27.31	19.07	21.59	41.29	က
92	303180	N92764	0.01	0.27	27.21	13.82	36.04	31.77	65
22	42415	R60981	0.12	3.11	26.71	9.46	66.35	5.30	<b>6</b>

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

m	က	<b>6</b>	60	က	က	၈	m	က	60	က	က	m	n	ന	ന	m	m	က	က	m	ო	m	က	က	က	m	က	ы	60	м	w	က	<b>6</b> 7	m	
36.33	13.87	25.64	14.11	12.48	26.60	10.88	32.50	9.98	26.43	11.55	49.72	24.62	22.42	9.30	40.27	20.16	14.13	10.73	16.88	34.04	31.84	32.84	5.91	33.48	9.40	26.82	12.25	6.49	5.36	15.00	22.27	5.13	26.44	5.25	
35.03	30.16	13.26	58.23	52.20	13.61	16.26	10.09	17.55	34.50	16.96	7.89	18.45	33.42	30.57	11.18	24.35	28.90	33.82	8.91	11.78	10.67	8.92	36.98	5.09	30.25	15.89	14.07	33.79	5.66	24.61	16.46	32.86	6.51	31.74	
7.36	33.86	38.89	5.18	12.08	34.18	46.38	30.26	44.37	8.11	38.20	5.07	18.18	5.36	20.94	7.9	10.20	10.45	7.52	25.95	5.64	8.16	7.84	6.72	10.47	9.38	5.81	21.48	6.61	33.97	5.14	5.11	5.64	9.65	5.44	
26.24	25.96	25.93	25.84	25.59	24.80	24.51	24.29	23.97	23.01	22.24	20.90	20.42	20.40	20.27	19.50	18.24	17,83	17.36	17.25	17.15	16.89	16.56	16.54	16.35	18.34	16.17	15.94	15.63	15.00	14.92	14.61	14.54	14.20	14.14	,
0.16	1.21	12.53	0.24	0.17	10.00	9.30	0.24	0.16	0.16	3.52	0.21	0.25	0.20	2.46	0.19	0.31	2.95	0.21	33.78	0.63	0.17	0.17	0.21	0.16	60.0	11.31	0.32	0.29	3.80	0.35	0.35	0.14	0.14	0:20	
0.01	9.05	0.48	0.0	0.01	0,40	0.38	0.01	000	600	0.16	0.0	0.0	0.01	0.12	0.01	200	0.17	0.0	95	000	0.0	0.0	00	10.0	10.0	0.70	0.02	0.02	0.25	0.02	0.02	0.0	0.0	20.0	
AA418903	H09818	H02294	R49555	PAGSAR	AAUU9840	AA778675	A A G 708 18	P20101	Means?	A A 0.2514.2	Wannas	H22824	N32542	H22563	R54036	A A RR3187	AA047441	AA701315	T49355	N47012	AA676998	WR1432	N28789	WB2263	H79466	W93717	T69270	AA418545	AA702335	H97140	R40434	AA437142	R39951	AA609189	
768056	46896	151184	38344	41505	430002	1049033	RRAZOD	2246	284744	265140	418094	51581	270975	51916	40010	1487161	488431	435099	67440	280155	454083	347586	269606	358990	235164	357373	R2869	767315	447416	250069	28298	757380	25915	1031446	
28	8	8	; ;	; p	; ;	\$ 8	, F	3 %	3 8	3 8	3 2	3 9	; 4	; 9	: ឡ	2 4	; ¥	φ 9	£ £	- <b>4</b>	. 64	? <b>2</b> 5	3 25	: 2	: :3	<b>3</b> 5	, K	. 95	22	, K	65	; ;;		B	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

ო	ო	ю	n	60	ы	n	ო	က	က	က	က	က	n	က	က	က	ຕ	m	"	es	67	• •••			<b>.</b>	es	rs •	m	6	en	က	es	က	က	က	
5.35	5.12	20.74	15.62	17.41	22.12	19.45	8.12	6.86	11.26	7.37	24.06	17.09	16.72	20.94	12.92	7.26	5.03	15.10	7.37	5.79	11.50	7.50		30.6	7.35	5.24	5.84	13.62	5.71	6.90	10.11	11.70	5.53	9.84	6.83	
5.86	29.82	9.76	11.85	15.99	8.92	12.04	9.42	17.85	21.72	7.81	7.58	13.80	8.33	7.88	8.10	13.18	5.77	8.56	10.28	40.79	11.73	7.79	7.40	13.22	11.24	12,44	8.66	8.45	8.47	12.47	7.21	9.17	13.78	10.99	8.33	
31.21	6.39	10.64	13.47	7.33	9,24	8.23	22.16	14.72	5.65	21.95	5.26	5.65	10.15	5.06	11.07	11.48	20.21	5.59	11.04	90 77	06.71	20.0	6.4/	5.48	9.36	10.06	12.99	5.38	13.21	7.38	9.14	5.30	69.9	5.02	9.78	
14.14	13.78	13.71	13.65	13.58	13.43	13.24	13.23	13.14	12.88	12.37	12.30	12.18	11.73	11.30	10.70	10.64	10.34	97.6	9.57		3.52	84.90 84.00	9.46	9.44	9.32	9.24	9.16	9.15	9.13	8.92	8.82	8.72	8.67	8.62	8.31	
1.25	0.39	15.63	16.52	0.17	0.13	8.63	191	2,3	210	35.	0.12	14.30	34.0	0.45	9 6	13.51	7 6.0	5 6	0.02 4	5 5	10.61	1.40	2.04	10.53	3.43	1.03	09'0	4.	1.40	0.11	0.05	4.90	12.90	3.90	157	:
0	0.03	4		100	00	0.65	3 5		3 6	5 6	3 6		2 2	3 3	3 6	10.4	1 2	* 6	S 6	70'0	1,12	0.15	0.22	1.12	0.37	0.11	0.0	0.16	0.15	0.0	100	55	97.5	97.0	9 0	;
097070	BAZEON	0002	A 608074	AA128617	AA016275	1402270	AARBEATE	20000	/1781U	N34042	4 6 6 4 4 6 6 9	A A 262244	***************************************	2005002	217761	LEGGA	100001	AA6/8295	AA041197	AA/01652	N47113	82106N	AA702684	N24042	AA045665	H04789	H77727	H46178	AA598817	W49672	Traced	AA197351	0.000000	AAE3000	DE0040	#1000U
32400	30850	407A70A	906308	50000	350250	745000	413009	200	24406	000447	777376	40.00	271000	1030130	70000	104483	3000	430926	3/6452	433567	280375	303089	448059	269300	489213	43733	234617	177884	897956	10467	78247	624627	707 707	#31.04 R26373	And A	307
G	3 2	5 4	3 8	8 6	5 8	8 6	8 6	? ;	: f	, t	2 ;	₹ ;	e ;	e ‡	: F	e 6	2 6	3 3	<b>5</b> 5	22	83	¥	82	98	87	8	68	8 6	8 8	<b>.</b> 6	: 8	3 3	5 2	3 8	8 8	ñ

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

ю	က	m	က	eo	ო	n	· •	. "	,	m (	<b>6</b> 3 (	י מ	ו מיז	m (	<b>.</b>	en	n	က	7	7	7	7	7	7	7	8	8	7	7	7	8	7	7	8	2	
8.51	5.12	9.05	7.97	11.50	7.97	976	7 7 7 9	) ; ;	DS: \	5.10 5	5.60	6.63	5.79	5.08	5.21	5.52	5.77	5.40 5.40	7.96	7.05	1917.92	5.67	6.10	9.57	9.33	7.73	1.35	5.84	146.14	3.45	105.59	12.54	13.76	7.41	99.88	
11.10	6.62	5.22	7.09	5.10	6.65		p 1	7.63	7.83	5.36	5.02	5.78	5.67	5.73	6.93	7.41	5.63	5.01	2866.06	2506.90	61.14	1831.96	1516.35	1171.33	944.19	841.07	674.68	302.09	140.40	249.63	150.46	239.88	229.57	203.18	95.72	!
5.27	11.86	8.56	77.7	20.00	7 30	D: -0	6.01	5,15	5.35	10.00	9.45	6.67	6.92	7.47	6.00	5.03	5.49	5.19	1.38	1,98	2.32	1.58	2.41	0.73	4.57	.0.	10.58	3.09	0.80	5.01	0.89	3.09	1.42	3.19	78.0	Š
8.30	7.87	7.61	7.61	7 55		ج <u>ا</u>	7.30	7.18	7.05	6.85	6.69	6.36	6.13	6.10	6.05	5.99	5.63	5.20	991.80	838.64	660.46	613.07	508.28	393.88	319.36	283.26	228.86	103.67	95.78	86.03	85.65	85.17	87. T.S.	7 52	27.33	64.00
7.83	0.54	69		20.0	2 (	3.90	0.12	18.03	3.05	0.72	2.07	47,65	7.33	16.66	29.59	0.07	1.94	0.12	9.92	83.8	9 16	. 6	00 K	200	بن در در در	2.83	1 70	2	. 63	2.66	. E	, r		20.0	- ;	ð
200	200	7, 0	2 1	S 5	0.02	0,53	0.02	2,51	0.43	0.10	0.31	7.49	1.20	2.73	8.89	500	75		5 6	5 6	5 6	5 6	5 5	5 6	5 6	5 5	5 6	5 6	5 6	, E	3 6	8 6	5 6	5 6	o i	0.08
AAAOKTEE	A A 4 E E D 3 B	200004	148084	AA608568	N30372	H29227	AA477400	AA663310	R06944	AA701645	528287	H02234	R10284	AA425302	A A 4 5 9 3 1 8	20227A	200001	20000	N32003	N/4000	20000	CSSOQN	N5/483	A4455003	N643/9	AA030444	#0667###	277000	M424300	AA074E03	AA40E34A	M4602214	/IRRCHW	11908/	AA8/6021	AA419486
	4222	177510	200402	069056	260035	. 52704	740620	853368	126650	433576	434749	454055	428047	773073	814108	27KB71	76037	30000	68704	800887	100000	4502/	217321	1/3040	290213	100010	/C110/	10000	077007	26603	200000	46/010	/7008/	347182	0581911	746152
ŝ	8 8	3	2	ğ	102	<b>50</b>	\$	105	404	3 5	2 5	2 5	3 5	2 =	: <b>;</b>	¥ ;	? ;	<b>*</b> 177	2 ;	91	<u>}</u>	81.	<del>1</del> 86	25	121	2 5	3 3	<u> </u>	3 ;	9 5	3 5	971	129	<u>දි</u>	<del>1</del> 3	132

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	8	7	7	64	~	7	~	~	~	7	7	~	~	~	~	7	~	~	7	7	7	7	~	7	~	~	7	~	7	7	84	7	7	7	
11.76	6.73	4.30	70.45	123.70	0.72	1.93	5.39	6.47	12.17	91.51	7.45	6.23	3.16	11.05	19.33	5.11	2.18	51.22	6.23	5.61	7.60	55.73	81.28	5.08	0.70	11.03	11.15	71.43	52.45	1.00	63.66	10,31	9.84	6.05	
168.10	173,62	170.43	4.69	51.92	88.50	154.39	150.40	1.37	139.36	56.58	128.30	120.65	121.33	109.53	99.20	105.34	105.49	52.03	89.38	0.12	81.20	34.54	7.97	76.38	9.84	70.25	69.04	1.36	27.83	72.00	11.75	66.04	62.71	68.60	
3.73	6.88	6.29	104.24	1.03	78.84	5.36	0.54	144.61	0.63	0.91	2.24	3.54	29'9	4.59	4.81	3.08	5.72	0.93	1.79	87.92	2.89	0.77	1.00	1.51	72.37	0.74	1.21	8.33	0.69	5.77	2.28	0.59	4.23	2.05	
61.20	60.41	60.34	59.79	58.88	56.02	53.90	52.11	50.82	50.72	49.67	45.99	43.48	43.37	41.72	41,11	38.18	37.79	34.73	32.46	31.22	30.56	30.35	30.08	27.65	27.63	27.34	27.13	27.04	26.99	26.26	25.90	25.65	25.59	25.57	ည
0.61	0.83	1.60	26.93	2.87	12.41	0.28	2.21	60.94	1.1	13.02	0.60	0.81	0.43	0.42	0.41	0.38	0.38	8.56	1.12	2.62	0.70	7.89	0.30	5.88	16.16	0.27	0.27	0.20	14.38	0.26	14.15	0.39	0.29	1.90	
00	0.0	9.03	0.45	0.05	0.22	0.01	9.0	1.20	0.02	0.26	0.01	0.02	0.01	0.0	0.0	0.01	0.0	0.25	9.03	90.0	0.02	0.26	0.0	0.21	0.58	0.0	0.01	0.0	0.53	0.01	0.55	0.02	0.01	0.07	
VA/04789	AA757754	AA461522	R23270	AA251784	W73144	AA086475	R40231	AA437224	R44741	AA504844	AA449120	AA855158	R37395	AA437126	AA810225	AA062985	AA857748	R91566	R59473	H29290	AA401434	AA485052	AA099369	R49439	R90934	R15740	AA460274	AA443624	H50655	AA425158	N47445	N52767	AA452139	R40018	
346942	395708	795858	131452	684879	344589	562811	30207	757435	33827	825740	785849	1389018	28498	757368	1367900	366209	1435339	196544	37823	49858	743189	815861	489656	38347	167205	53039	796549	771294	194318	769000	280699	283379	787856	27605	
133	<u>\$</u>	55	136 85	137	138	139	5	<del>1</del>	142	<u> </u>	4	145	3	14	148	149	8	151	152	153	\$	55	55	167	85	159	6	161	162	163	<b>3</b> 5	165	166	167	

· Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	8	<b>6</b> 1	2			7	7		7	7	. 7	74	6	7	8	73	2	2	2	7	77	71	7	8	7	7	7	2	<b>7</b>	7	2	2
16.63	5.46	8.05	11.62	31.35	45.50	11.25	62.76	43.13	3.46	48.25	25.34	31.76	11.14	53.51	54.20	23.77	3.08	47.62	17.48	28.27	20.17	50.17	45.41	25.26	49.74	46.16	1.08	43.06	5.54	10.71	6.22	5.54	16.22	8.77
26.22	67.88	64.40	60.05	41.17	26.82	60.48	9.54	28.65	59.71	21.91	42.67	30.37	51.95	9.53	10.05	1.50	23.72	10.88	42.76	30.54	38.72	7.58	10.31	30.81	1.00	1.93	14.31	10.12	47.06	38.83	45.91	45.44	34.15	40.61
													3.82																		٠			
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500													0.17 3.74														••	•						
	0 F640F04		NISCIA C										P4463450 C																					•
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,	9 5	ĝ (	0. i	<u> </u>	172	173	<b>1</b> 5	175	176	Ē į	0/1	n 6	<u> </u>	<u>.</u>	70.	2 5	Ē	5 <b>5</b>	487	, <del>,</del>	3 2	9 9	<u></u>	ç Ç	£	3	195	85	197	801	90	200	3 5	:

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	7	7	2	2	~	7	. 7	7	8	7	7	8	7	7	~	7	7	7	~	8	2	7	7	8	7	7	7	~	7	7	8	
5.21	12.21	34.18	7.22	5.65	6.69	28.36	25.15	19.22	15.13	18.34	6.87	15.64	23.25	28.45	28.16	5.10	27.66	4.48	12.18	18.27	29.19	18.18	7.18	26.50	7.37	25.70	28.39	2.23	3.21	5.10	0.58	7.04	26.39	20.98	
42.11	37.48	1.51	39.78	40.90	36.93	16.52	21.09	22.78	28.43	26.10	0.27	3.02	18.93	4.19	8.24	36.28	11.04	18.97	26.44	21.58	8.82	18.08	29.16	1.54	0.78	12.28	8.17	29.01	28.51	31.49	14.55	29.00	0.39	14.19	
4.26	1.87	14.65	2.86	2.84	3.94	2.63	0.69	4.69	1.53	0.42	36.64	24.69	1.00	10.07	5.00	0.88	2.99	17.85	2,42	990.	2.08	3.39	3.08	10.97	30.80	0.65	1.86	7.14	6.31	1.33	22.57	1.18	10.40	1.96	
17.19	17.19	16.78	16.62	16.47	15.85	15.84	15.64	15.57	15.03	14.95	14.60	14.41	14.40	14.24	14.13	14.09	13.89	13.76	13.68	13.50	13.36	13.22	13.14	13.00	12.97	12.88	12.81	12.79	12.67	12.64	12.56	12.41	12.39	12.38	
0.86	1.06	0.11	0.29	1.14	0.40	2.94	5,98	0.30	3.91	3.94	25.0	<b>7</b> .	0.14	3.51	17.58	2.19	4.17	0.13	1.31	3.95	8.41	0.26	1.05	0.08	0.36	0.13	8.32	0.71	0.25	1.99	0.13	4.99	9,44	0.12	
0.05	90.0	0.01	0.02	0.07	0.03	0.19	0.38	0.02	0.26	0.26	90.0	0.11	0.01	0.25	1.24	0.16	0.30	0.0	0.10	0.29	0.63	0.02	0.08	0.01	0.03	0.0	0.65	90.0	0.02	0.16	0.01	0.40	0.76	0.01	
AA609304	AA857163	AA481481	N26559	N51297	AA410437	AA504838	AA417713	AA045175	AA463445	H18932	AA035745	AA476543	AA434388	AA883800	AA774665	AA699601	W90705	R56149	R37738	AA427719	H15549	AA491227	AA883181	N52193	AA035730	AA017379	AA911236	R43915	AA621355	AA070435	AA453618	AA437212	AA620697	AA022496	
1031588	1410444	756629	266366	283070	753411	825726	746232	487766	811766	51599	364865	785293	770860	1461609	856289	436402	418004	41406	26808	771004	49499	824074	1467166	284288	364844	361122	1526789	33066	1055121	530954	795446	757428	1049185	384352	
203	707	206	506	202	208	503	210	211	212	213	214	215	216	217	218	219	22	Z	22	82	727	225	228	727	228	229	230	23.	232	233	8	235	238	237	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

η (	7 (	η (	71	7 (	7 (	<b>v</b> (	70	7	N (	<b>(N</b> )	7 (	7 (	<b>74</b> (	71	<b>74</b>	7	7	N	7	7	7	8	8	71	8		1 0		<b>v</b> (	<b>7</b>	7	<b>~</b>	8	7	7	
11.09	<del>-</del> 8	17.55	23.17	20.10	12.88	18.24	19.11	14.01	12.09	3.40	20.99	24.11	16.34	7.83	13.80	4. 2.	16.62	10.20	12.02	20.10	8.36	5.22	21 77	4	87.8	2 4	13.60	23.70	15.64	0.48	1.65	17.42	21.42	5.78	1.12	
23.87	29.43	15.47	4.23	14.95	18.70	17.13	13.18	20.54	19.31	20.61	10.18	5.70	16.85	22.78	17.96	9.83	15.37	20.21	15.78	10.22	1.00	22.31	9	6.6	2 2	21.65	14.48	6.31	14.23	7.21	17.15	11.85	6.06	20.48	20.18	
211	6.29	3.28	8.77	1.00	4. 4.	2.08	3.09	0.58	3.60	10.87	3.65	4.58	0.33	2.56	0.92	17.71	0.51	2.09	4.64	2.05	22.91	7 63	r i		06.12	2.67	0.88	0.86	0.77	22.87	11,51	0.93	2.67	3.74	198	<u>;</u>
12.36	12.24	12.10	12.06	12.02	11.98	11.82	11.80	11.71	11.67	11.63	11.61	11.46	11.17	11.06	10.90	10.87	10.84	10.83	10.81	40.79	10.76	0.01	10.00	10.50	10.45	10.35	10.32	10.31	10.21	10.19	10.11	10.07	10.05	60 01	0 02	i. 6. 6.
3.72	0.12	0.31	16.58	1.69	16.82	23.53	3.89	0.18	11.24	0.20	27.64	0.11	0.34	0.95	3.90	5.68	6.12	0.34			7.6	. i	5.93	8.09	12.13	9.4	0.87	4.21	5.55	9.47	0.12	2.53	15.05	76.0	2	767
0:30	0.01	0.03	1.38	0.14	1.40	1.99	0.33	0 02	96.0	0.02	2.38	0.0	0.03	0.09	0.36	0.52	0.57		3 5		8 3 3	0.0	0.56	0.77	1.18	9	0.08	0.41	0.54	0.83	0.0	0.25	3 5	3 6	<b>5</b>	0.24
H10713	T57927	AA778717	A475575	R92201	AA706901	AAATT22T	HOODE	A 4 E 1 4 8 8	D44818	W03523	AA026587	AA179161	AA160670	AA427737	H48070	AA460389	AA410474	1	N/4131	AA461513	N30751	AA680136	AA417920	AA489073	AA454949	N93721	AA281784	H54451	N49389	AA412049	AA158035	2002	20025	A4478038	AA598787	AA626167
47764	71871	4049079	705368	195845	451640	720450	JEARI	504467	01,00	357138	366071	600278	507738	770854	40333	705746	753/78	133420	23641	782840	257955	433155	767403	824802	814769	307138	712401	203184	277621	220053	500640	407664	18/02/	/4045/	898073	745248
ž	3 8	600	} ;	242	. 5	? ?	į	2 6	2 2	<b>3</b> 7	2 0	, y	3 4	25.2	7 5	3 3	g :	65	529	727	258	259	260	261	797	263	284	265	266	757	3 8	8 6	807	270	27	<b>13</b>

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

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9.80 5.59 9.79 3.40
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AA682533 AA620528
391987 490387 431242 951305
- 1, 1 1 0

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

8	~	7	7	7	7	7	7	7	7	7	2	7	7	7	7	7	~4	7	7	7	7	7	8	7	7	6	7	2	2	7	7	7	7	7	
14.67	5.38	9.46	1.10	16.60	5.40	26,1	8.45	14.43	8.30	18.10	15,48	7.50	14,35	18.86	3.90	17.99	11.27	12.36	17.21	6.37	14.40	16.24	19.27	2.86	13.09	11.28	18.07	7.71	14.93	15.23	9.43	5.36	13.11	13.46	
10.87	20.24	15.22	13.45	0:30	20.84	2.96	17.16	2.08	17.44	6.64	6.65	16.58	1.00	0.52	14.21	6.50	12.46	11.51	6.16	13.73	6.65	6.27	0.41	13.18	1.67	11.14	0.56	13.64	5.10	3.98	13.54	18.34	7.53	8.34	
1.65	1.54	2.44	12.54	10.05	0.65	11.72	0.92	9.92	0.36	4.5	3.86	1.87	10.33	6.22	7.42	0.78	1.45	1.30	1.81	4.95	3.89	2.41	5.22	8.74	9.98	2.23	5.83	3.04	4.35	5.06	2.7	0.38	3.42	2.25	
9.06	9.06	9.04	9.03	8.98	8.96	8.87	8.84	8.81	8.70	8.69	8.66	8.65	8.56	8.53	8.51	8.42	8.39	8.39	8.39	8.35	8.32	8.31	8.30	8.29	8.25	8.21	8.15	8.13	8.13	8.09	8.08	8.03	8.02	8.02	
3.21	3.62	10.62	11.71	17.81	0.21	5.97	3.32	2.18	39,21	0.80	48.92	34.28	0.09	5.11	0.14	6.98	9.51	3.36	0.08	0.19	4.39	13.45	10.75	1.09	0.15	1.49	0.14	0.25	21.34	10.44	6.11	0.22	0.28	90.0	
0.35	0.40	17.	1.30	1.98	0.02	0.67	0.38	0.25	4.51	0.09	5.65	3.96	0.0	0.60	0.02	0.83	1.13	0.40	0.0	0.02	0.53	<u> 5</u>	8.	0.13	0.02	0.18	20:0	0.03	2.63	1.29	0.78	0.03	0.03	0.0	
N26515	AA192527	AA488175	AA172056	AA620485	AA416970	W68220	H41203	H16772	AA628999	AA778646	H95638	AA775863	AA486998	AA682321	T52311	N80764	W47364	AA504265	AA036974	AA485265	AA455786	N74617	AA488324	R61372	N50806	AA058578	AA131239	W96216	R41329	H54752	AA699427	AA457576	AA705825	H94944	
266259	627541	877632	594758	951241	730036	342640	175533	50227	884511	1048995	242952	878544	897546	462926	71977	300615	324717	825416	484535	840150	809557	298429	842968	37880	280882	380884	503581	358599	30148	203275	433253	838776	1292182	230261	
308	808	340	3.5	312	313	314	34.5	318	317	318	319	320	32	322	323	324	325	326	327	328	329	330	331	332	333	134	92	338	337	338	330	<b>2</b>	7	3 3	!

2	7	8	2	7	7	2	7	7	7	62	7	7	7	~	~	7	7	~	8	7	7	7	7	~	7	7	<b>7</b> 1	8	8	2	7	7	7	8	
8.40	8.68	6.01	6.49	10.12	4,46	15.20	40.00	11.39	10.84	9.28	13.28	<del>.</del> .	11.71	9.03	6.8 29.	2.00	5.66	9.32	15.79	7.15	15.99	88.6	8.23	15.83	9.18	5.28	12.04	8.29	4.93	12.46	12.18	8.02	14.40	10.67	
14.68	13.94	15.21	11.97	9.24	7.24	7.43	8.93	4.17	7.66	12.63	3.58	14.11	8.00	11.89	14.72	8.59	16.02	12.70	5.74	4.96	4.00	12.36	10.88	5.39	0.92	4.41	8.97	11.98	6.94	4.68	6.84	13.66	0.63	0.08	
0.92	1.06	227	4.86	3.94	11.54	75.0	4.23	7.55	4.60	1.18	6.23	7.96	3.35	5.09	1.41	12.36	123	0.83	121	10.53	5.64	0.35	3.47	125	12.26	12.64	1.30	0.99	10.35	5.03	3.00	9.34	6.83	11.07	
8.00	7.89	7.83	11.1	77.7	7.75	7.73	7.72	7.70	7.70	7.70	7.70	7.69	7.68	7.67	7.66	7.65	7.64	7.61	7.60	7.55	7.54	7.53	7.53	7.40	7.46	7.44	7.44	7.41	7.41	7.39	7.34	7.34	7.29	72.7	
28.81	2.29	5.20	2.89	0.60	5.81	0.14	0.35	24.59	13.42	2.05	1.68	0.08	261.61	0.53	0.25	1.03	18.06	0.08	5.46	0.04	0.08	7.28	20.00	1.18	0.08	0.20	6.14	0.31	21.88	23.90	57.27	9.9	28.27	15.11	
	2 2	<b>9</b>	0.37	0.08	0.75	0.02	0.05	3.19	1.74	0.27	020	0.0	8	0.07	0.03	0.13	2.36	0.01	22.0	0.01	0.0	76'0	0.75	0.16	0.04	0.03	0.83	90.0	2.95	3.23	7.80	55.	3.88	2.08	
1061768	C0/16U	DE0775	H20807	A4625764	0.02010	AA158795	AA906257	196718	155607	H57082	W15305	A 4 4 6 3 9 R 2	AA676460	H10231	H10068	AA780365	W72666	195804	AA291773	AA070331	AA702802	R53455	AA434403	AA633835	AA485749	T61122	R12473	R49644	AA701455	N93924	AA488367	N82936	T66816	AA706968	
403803	701635	250107	10035	745393	782759	502444	1505919	121251	73596	204644	3224B1	810581	882510	46927	46740	1035664	345743	120695	725340	530820	448032	39833	770879	868169	811168	83666	128243	37538	435076	309288	843163	278729	66317	451907	
5	} ;	<b>*</b>	3	3	ξ	2	350	×	352	353	3	3,5	<b>3</b>	357	358	359	360	361	362	363	364	365	366	367	88	369	370	37.	372	373	374	375	376	377	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line; Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	8	2	7	7	7	. ~	, ,			• •	• •	<b>4</b> r	, r	• •	<b>4</b> (	7 (	,	74	2 •	7 (	7	7	N (	N (	٧ (	N (	71 (	7	7	7	7	8	2	
10.22	12.41	6.82	6.11	10.54	6.43	, K	) O	44.97	3.0	10.38	60.01	9.5	40.7	7.62	5.32	40.0	6.81	3.65	6.50	11.87	9.20	14.49	11.35	9.27	10.38	1.16	3.14	9.86	5.69	8.65	12.53	8.19	9.34	11.61	5.09	
9.92	2.06	13.06	12.65	7.03	43 AF	6.5	11.90	00.0	19.7	8.89 10.00	6.35	8.75	0.03	11.28	2.34	11,35	12.50	5.87	9.95	7.50	7.07	5.67	6.52	9.22	8.38	5.14	11,26	4.41	10.31	9.72	1.10	1.51	8.48	7.15	13.88	
1.57	7.23	1.81	200		- ;	0.10	4.30 9.50	3.01	2.72	2.25	4.40	0.87	13.51	2.47	13.68	1.14	2.00	11.72	4.65	1.68	4.74	0.85	3.01	2.37	2.09	14.50	6.40	6.49	4.76	2.34	7.04	10.95	2.73	1.77	1,52	
7.24	7.23	7.23		S : 1	57.7	7.22	7.22	7.20	7.20	7.17	7.15	7.14	7.13	7.12	7.11	7.11	7.11	7.08	7.03	7.02	7.00	7.00	6.96	6.95	6.95	6.93	6.93	6.92	6.92	6.90	6.89	6.88	6.85	6.84	6.83	12
0.35	7 20	17	= :	50. 1 50. 1	5.80	0.31	33.83	6.75	4.04 40.4	15.96	0.10	0.11	105.24	14.44	0.62	1.51	95.95	14.78	0.08	2.01	100.32	0.28	4.0	0.68	36.26	0.07	0.39	3.69	7.46	3.78	90.0	2.74	4.75	0.83	0.25	
300	3 6	5 6	9.70	1.24	0.80	9.0	4.68	0.94	0.64	2.23	0.01	0.02	14.77	203	60.0	0.21	13.50	2.09	50	520	14.33	80	90'0	0.10	5.22	10.0	90.0	0.63	1,08	0.55	5	0.40	990	0.12	90	
992CC14	N2270	K43308	H19234	AA135809	T62072	AA011182	H97488	AA460952	AA45521	AA504201	AA456611	AA677655	AA894557	AA455237	AA449118	R96520	NEGISB	N45236	247503	K/4503	AINTERED	WTOKK	MARTOR	TARATA	A A 4 R 5 4 5 3	T58146	RAS048	N20305	AA465238	A A 886710	A A 660273	AA452278	AA420436	AIM5250	AA398408	
	2000	32756	50904	\$65734	85670	359722	251936	796114	809828	825282	809567	897271	1416782	814779	785845	100544	278534	282171	44006	2,7050	000/47	345330	44600	#1000 #1000	044067	70254	20773	264502	814225	1600463	900106	707010	000.00	1844287	728704	
	378	379	380	381	382	383	38	385	386	387	88	388	067	5	392	, ot	3 8	30.		999	A 6	0 0	n (	3 5	·	<b>7</b>	3 3	į	5 5	3 5	) (	ş ç	3 5	2 ;	# <del>*</del> *	<u>.</u>

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

204560         H68000         0.41         2.79         6.80         1.22         12.71         6.46         2.2           204560         H68000         0.01         0.03         6.79         6.79         6.77	131316	R23055	1.19	8.14	6.82	3.20	5.03	12.23	
AA266231         0,01         0,08         6.79         4,75         10.24           H71230         0,05         0,34         6,79         6,29         4.78           H08595         1,32         8,93         6,79         6,29         4.78           H08595         1,32         8,93         6,79         6,29         4.78           AA454562         0,24         1,65         6,77         12.08         7.25           AA454567         1,50         10,08         6,77         12.08         7.25           AA465697         0,24         0,68         6,72         7.84         6.89         4.69           AA465693         0,54         0,69         6,72         7.84         6.81         1.62           AA465693         0,54         0,62         6,65         0,66         3.37         9.66           AA436899         0,70         4,67         6,65         7.44         4.89         4.47           AA436899         0,70         4,67         6,65         7.39         4.47         4.47           AA43689         0,70         4,67         6,66         3.37         9.66         4.47           AA48689         0,70	8	H56903	0.41	2.79	6.80	1.22	12.71	6.48	7
H71201         0.05         0.34         6.79         6.29         4.78           H08595         1.32         8.93         6.79         6.37         9.34           AA010085         4.09         27.70         6.77         5.57         3.78           AA454562         0.24         1.65         6.77         1.208         7.25           AA454567         1.50         10.08         6.72         1.82         4.68           AA454567         1.50         10.08         6.71         2.08         1.82           AA455872         0.54         3.61         6.72         8.09         1.073           AA559032         1.11         7.44         6.71         2.08         4.47           AA458942         0.70         4.67         6.65         3.37         9.66           AA45895         0.70         4.67         6.65         7.37         9.68           AA45896         0.01         0.07         6.65         7.87         10.05           AA44800         0.01         0.07         6.65         7.37         10.05           AA48400         0.01         0.07         6.64         11.27         0.58           H5572B </td <th>9</th> <td>AA256231</td> <td>0.0</td> <td>90:0</td> <td>6.79</td> <td>4.75</td> <td>10.24</td> <td>5.38</td> <td>7</td>	9	AA256231	0.0	90:0	6.79	4.75	10.24	5.38	7
AAA454562         1,32         8,93         6,79         6,37         9,34           AAA454562         0,24         1,65         6,77         1,208         7,25           AAA454567         1,50         1,008         6,77         1,208         7,25           AAA45457         1,50         1,008         6,77         1,208         7,25           AAA45457         1,50         1,008         6,77         1,208         7,25           AAA45457         0,04         0,30         6,72         7,84         4,68           AAA539949         1,11         7,44         6,71         2,08         1,073           AAA46593         9,85         6,56         6,65         7,13         4,39         9,66           AAA46693         0,70         4,67         6,65         7,13         4,47         4,47           AAA46693         0,70         4,67         6,65         7,13         4,39         9,66         9,66         9,56         4,47           AA44600         0,01         0,07         6,65         7,13         4,37         4,47           AA448400         0,01         0,07         6,65         7,26         4,47 <t< td=""><th>8</th><td>H71230</td><td>0.05</td><td>9. 24.</td><td>6.79</td><td>9.29</td><td>4.78</td><td>6.30</td><td>7</td></t<>	8	H71230	0.05	9. 24.	6.79	9.29	4.78	6.30	7
AA010055         4.09         27.70         6.77         5.57         37.8           AA454567         0.24         1.65         6.77         12.08         7.25           AA454597         1.50         10.08         6.74         9.56         1.62           AA454597         1.50         10.08         6.72         7.84         4.68           AA46597         1.50         10.08         6.72         7.84         4.68           AA46593         0.54         3.61         6.72         7.84         4.68           AA598949         1.11         7.44         6.71         2.08         10.73           AA458902         2.85         65.62         6.66         3.37         9.66           AA458902         1.37         7.73         6.67         7.33         4.39           AA44600         0.17         7.75         6.64         1.12         0.66           AA44400         0.01         0.07         6.65         7.44         4.75           AA44400         0.01         0.07         6.65         7.85         4.47           AA44400         0.01         0.07         6.59         7.52         7.44           AA48605	<u> </u>	H08595	1.32	8.93	6.79	6.37	9.34	4.64	6
AA454582         0.24         1.65         6.77         12.08         7.25           AA454587         1.50         10.08         6.74         9.56         1.68           H17139         0.04         0.30         6.72         7.84         4.68           AA609723         0.54         3.61         6.71         2.08         1.63           AA458949         1.11         7.44         6.71         2.08         10.73           AA458902         2.82         6.56.2         6.66         3.71         4.89           AA458902         2.82         18.79         6.65         7.13         4.39           AA465893         0.70         4.67         6.65         7.13         4.39           AA44600         0.71         7.75         6.64         1.12         0.56           AA44400         0.01         0.77         6.65         7.44         7.59           AA44400         0.01         0.77         6.65         7.44         7.59           AA44400         0.01         0.77         6.65         7.44         7.59           AA44400         0.01         0.77         6.59         7.24         4.74           AA4800 <th>6</th> <td>AA010065</td> <td>4.09</td> <td>27.70</td> <td>6.77</td> <td>5.57</td> <td>3.78</td> <td>10.97</td> <td>7</td>	6	AA010065	4.09	27.70	6.77	5.57	3.78	10.97	7
AAAGSGS         1.50         10.08         6.74         9.56         1.82           H17139         0.04         0.30         6.72         7.84         4.68           AAA039049         1.11         7.44         6.71         2.08         10.73           AAA58902         2.82         3.61         6.72         8.02         8.01           AAA58902         2.82         18.79         6.65         6.66         3.37         9.66           AAA456895         0.70         4.67         6.65         7.13         4.47           AAA456895         0.70         4.67         6.65         7.13         4.47           AAA44800         0.70         4.67         6.65         7.13         4.47           AAA44800         0.01         0.07         6.64         8.09         3.71           H55728         9.61         1.127         0.58         4.47           AA44800         0.01         0.07         6.62         7.44         7.44           AA44800         0.01         0.07         6.62         7.87         4.47         4.47           AA44806         0.01         0.07         6.62         7.84         4.47         4.47 </td <th>2</th> <td>AA454562</td> <td>0.24</td> <td>1.65</td> <td>6.77</td> <td>12.08</td> <td>7.25</td> <td>0.98</td> <td>74</td>	2	AA454562	0.24	1.65	6.77	12.08	7.25	0.98	74
H17139         0.04         0.30         6.72         7.84         4.68           AAB09723         0.54         3.61         6.72         8.02         8.01           AAB09723         0.54         3.61         6.72         8.02         8.01           AAB09723         1.11         7.44         6.71         2.08         10.73           AAB09022         2.82         18.79         6.65         3.37         9.66           AAB09022         2.82         18.79         6.65         7.13         4.39           AAB09022         2.82         18.79         6.65         7.13         4.39           AAB09022         1.77         7.75         6.64         10.73         4.47           AAA44000         0.01         0.07         6.62         7.87         10.05           AAA44000         0.01         0.07         6.62         7.87         10.05           AAA44000         0.01         0.07         6.64         8.09         3.71           AAA08400         0.01         0.07         6.62         7.87         10.05           AAA08400         0.01         0.07         6.64         8.09         3.71           AA	82	AA454597	1.50	10.08	6.74	9.56	1.82	8.83	7
AAG99723         0.54         3,61         6,72         8,02         8,01           AAJ39849         1,11         7,44         6,71         2,08         10,73           AA458902         1,11         7,44         6,71         2,08         10,73           AA59802         2,82         18,79         6,65         3,37         9,65           AA59802         2,82         18,79         6,65         7,13         4,39           AA45689         0,70         4,67         6,65         7,13         4,47           AA45689         0,70         4,67         6,64         11,27         0,58           AA448400         0,01         0,07         6,62         7,97         0,68           AA448400         0,01         0,07         6,62         7,97         0,08           AA448400         0,01         0,07         6,59         7,24         7,59           AA48040	83	H17139	90.0	0.30	6.72	7.84	4.68	7.65	~
AA399949         1.11         7.44         6.71         2.08         10.73           AA46593         9.85         65.62         6.66         3.37         9.66           AA456902         2.82         18.79         6.65         7.13         4.39           AA456902         2.82         18.79         6.65         7.13         4.39           AA456902         0.70         4.67         6.65         7.13         4.39           AA456902         0.70         4.67         6.65         7.13         4.39           AA45603         0.70         4.67         6.64         11.27         0.58           AA448400         0.01         0.07         6.62         7.87         4.47           AA448400         0.01         0.07         6.62         7.87         4.47           AA448400         0.01         0.07         6.59         7.44         4.47           AA448400         0.01         0.07         6.59         7.24         7.59           AA448400         0.02         0.01         6.57         6.57         7.44           AA439851         0.02         0.12         6.54         4.93         6.59           AA430	883	AA609723	45.0	3.61	6.72	8.02	8.01	4:14	8
AA456593         9.85         65.62         66.6         3.37         9.65           AA599022         2.82         18.79         6.65         7.13         4.39           AA456895         0.70         4.67         6.65         7.13         4.39           AA4456895         0.70         4.67         6.65         7.13         4.39           AA445690         0.17         7.75         6.64         11.27         0.58           AA442790         0.17         7.75         6.64         11.27         0.58           AA442400         0.01         0.07         6.62         7.87         10.05           AA448400         0.01         0.07         6.59         7.44         9.16           AA448420         0.01         0.07         6.59         7.44         7.59           AA484202         0.01         0.07         6.59         7.52         7.44           AA894222         0.01         0.02         6.51         6.51         7.44         7.59           H05645         0.02         0.12         6.53         6.54         6.57         7.44         7.59           AA43087         0.04         0.24         6.53         4.	677	AA398949	=	7.44	6.71	2.08	10.73	7.33	7
AASSGOS         2.82         18.79         6.65         7.13         4.39           AAA456895         0.70         4.67         6.65         7.85         4.47           AAA42790         1.17         7.75         6.64         11.27         0.58           AAA48400         0.01         0.07         6.62         7.97         10.05           AA48400         0.01         0.07         6.62         7.97         10.05           AAA88452         0.01         0.07         6.62         7.97         10.05           AAA88452         0.01         0.07         6.62         7.87         10.05           AAA38851         0.02         0.12         6.57         2.44         7.59           H05645         0.06         0.22         6.57         2.44         7.59           H05645         0.06         0.42         6.54         1.08         5.69           H05645	24B	AA465593	9.85	65.62	99'9	3.37	9.66	6.95	7
AA456895         0.70         4.67         6.65         7.85         4.47           AA426895         0.70         4.67         6.64         11.27         0.58           AA424790         1.17         7.75         6.64         1.127         0.58           AA448400         0.01         0.07         6.62         7.97         10.05           AA48400         0.01         0.07         6.62         7.97         10.05           AA48400         0.01         0.07         6.59         7.52         7.44           AA484522         0.01         0.07         6.59         7.52         7.44           AA039851         0.02         0.12         6.57         2.44         7.59           HD5645         0.06         0.42         6.57         2.44         7.59           HD5645         0.06         0.42         6.57         2.44         7.59           HD5646         0.06         0.42         6.57         2.44         7.59           HD5647         0.11         6.53         5.27         9.69           HB447         6.53         5.27         9.69           AA428166         0.02         0.12         6.46	445	AA599092	2.82	18.79	6.65	7.13	4.39	8.44	8
AA424790         1,17         7,75         6.64         11,27         0.68           H5203         0.68         4,52         6.64         8.09         3.71           H5203         0.68         4,52         6.64         8.09         3.71           AA448400         0.01         0.07         6.62         7.97         10.05           AA884622         0.01         0.07         6.59         7.52         7.44           AA038851         0.02         0.12         6.57         2.44         7.59           H05645         0.06         0.42         6.54         10.89         5.69           H20847         1.12         7.31         6.53         5.27         9.69           H20845         0.06         0.42         6.54         10.89         5.69           H20847         1.12         7.31         6.53         5.27         9.69           H21943         1.01         6.55         6.50         2.35         11.59           AA428166         0.02         0.24         6.46         2.50         9.29           AA428168         4.30         27.63         6.46         2.50         9.29           AA452542	149	AA456895	0.70	4.67	6.65	7.85	4.47	7.62	7
H59203         0.68         4.52         6.64         8.09         3.71           AA448400         0.01         0.07         6.62         7.07         10.05           T55728         9.61         63.60         6.62         7.07         10.05           AA884622         0.01         0.07         6.59         7.52         7.44           AA039851         0.02         0.12         6.57         2.44         7.59           HD5645         0.06         0.42         6.54         10.39         5.69           HD645         0.06         0.42         6.54         10.39         5.69           H20847         1.12         7.31         6.53         5.27         9.69           N48075         0.41         2.64         6.52         4.93         8.97           R43328         1.01         6.55         6.50         2.35         11.59           AA428186         0.02         0.12         6.46         2.35         11.59           AA428186         0.02         0.12         6.46         2.50         9.29           AA428186         0.02         0.12         6.46         2.50         9.29           AA428186	172	AA424790	1.17	7.75	6.64	11.27	0.58	8.09	8
AA448400         0.01         0.07         6.62         7.87         10.05           T55728         9.61         63.60         6.62         1.16         9.16           AA884622         0.01         0.07         6.59         7.52         7.44           AA039851         0.02         0.12         6.57         2.44         7.59           HDS645         0.06         0.42         6.54         10.89         5.69           HZ0847         1.12         7.31         6.53         5.27         9.69           HZ0847         1.12         7.31         6.52         4.93         8.97           NA8075         0.41         2.64         6.52         4.93         8.97           R43288         1.01         6.55         6.51         1.47         11.33           AA42161         0.04         0.24         6.51         1.47         11.33           AA42284         0.02         0.12         6.46         2.50         2.55         9.69           AA42542         0.07         0.12         6.46         2.50         2.25         9.13           AA445684         1.05         6.74         6.41         2.05         9.13     <	214	H59203	0.68	4.52	6.64	8.09	3.71	8.12	7
T55728         9.61         63.60         6.62         1.16         9.16           AA884622         0.01         0.07         6.59         7.52         7.44           AA039851         0.02         0.12         6.57         2.44         7.59           H05645         0.06         0.42         6.54         10.89         5.69           H20847         1.12         7.31         6.53         5.27         9.69           H20847         1.12         7.31         6.53         5.27         9.69           N48075         0.41         2.64         6.52         4.93         8.97           N48075         0.41         2.64         6.52         4.93         8.97           N48075         0.41         6.55         6.50         2.35         11.59           AA95326         1.01         6.55         6.50         2.35         11.59           AA428166         0.02         0.12         6.46         2.50         9.29           AA428166         0.02         0.12         6.46         2.50         9.29           AA428166         0.02         0.12         6.46         2.50         9.13           AA428166	362	AA448400	0.01	0.0	6.62	7.97	10.05	1.85	74
AA038651         0.01         0.07         6.59         7.52         7.44           AA038651         0.02         0.12         6.57         2.44         7.59           HD5645         0.06         0.42         6.54         10.89         5.69           HD6847         1.12         7.31         6.53         5.27         9.69           NA8075         0.41         2.64         6.52         4.93         6.97           T51617         0.04         0.24         6.51         1.47         11.33           R43528         1.01         6.55         6.50         2.35         11.59           AA953357         1.34         8.71         6.49         10.79         2.65           AA428186         0.02         0.12         6.46         2.50         9.29           AA428186         0.02         0.12         6.46         2.50         9.29           AA42818         0.02         0.12         6.46         2.50         9.13           AA42868         1.30         27.63         6.43         1.36         9.13           AA45542         0.07         0.45         6.41         2.86         6.70           AA45546	538	T55728	9.61	63.60	6.62	1.16	9.16	9.54	ત
AA039851         0.02         0.12         6.57         2.44         7.59           H05645         0.06         0.42         6.54         10.89         5.69           H05645         0.06         0.42         6.53         5.27         9.69           NA8075         0.41         2.64         6.52         4.83         8.97           T51617         0.04         0.24         6.51         1.47         11.33           R43328         1.01         6.55         6.50         2.35         11.59           AA953357         1.34         8.71         6.49         10.79         2.65           AA428186         0.02         0.12         6.46         2.50         9.29           AA42868         1.13         7.26         6.45         3.07         8.32           AA45542         0.07         0.45         6.43         1.38         9.13           AA45542         0.07         0.45         6.41         2.05         11.14           AA45542         0.07         0.45         6.41         2.05         11.14           AA45542         0.07         0.45         6.41         2.05         6.70           AA757564	8597	AA884622	0.0	0.07	6.59	7.52	7.44	4.80	8
H05645         0.06         0.42         6.54         10.89         5.69           H20847         1.12         7.31         6.53         5.27         9.69           NA8075         0.41         2.64         6.52         4.93         8.97           NA8075         0.41         2.64         6.52         4.93         8.97           T51617         0.04         0.24         6.51         1.47         11.33           R43228         1.01         6.55         6.50         2.35         11.59           AA953357         1.34         8.71         6.49         10.79         2.65           H21943         1.30         8.44         6.48         4.28         5.16           AA428186         0.02         0.12         6.46         2.50         9.29           AA428186         0.02         0.12         6.45         3.07         8.32           AA428788         4.30         27.63         6.43         1.38         9.13           AA448644         1.05         6.74         6.41         2.85         6.70           AA448644         1.05         6.74         5.69         11.14           AA757584         0.13	827	AA039851	0.02	0.12	6.57	2.44	7.59	9.68	7
H20847 1.12 7.31 6.53 5.27 9.69  N48075 0.41 2.64 6.52 4.83 6.97  T51617 0.04 0.24 6.51 1.47 11.33  AA953357 1.34 8.71 6.49 10.79 2.65  AA042816 0.02 0.12 6.46 2.50 9.29  AA042816 0.02 0.12 6.46 2.50 9.29  AA045508 1.13 7.26 6.45 2.50 9.29  AA034115 0.26 1.70 6.43 1.38 9.13  AA448664 1.05 6.45 6.42 2.05 11.14  AA45542 0.07 0.45 6.41 2.85 6.70  AA75784 0.13 0.82 6.41 5.00 4.97  H16910 0.03 0.21 6.41 5.69 12.25	299	H05645	90.0	0.42	6.54	10.89	5.69	3.05	~
N48075         0.41         2.64         6.52         4.93         8.97           T51617         0.04         0.24         6.51         1.47         11.33           R43328         1.01         6.55         6.50         2.35         11.59           AA953357         1.34         8.71         6.49         10.79         2.65           H21943         1.30         8.44         6.48         4.26         5.16           AA428186         0.02         0.12         6.46         2.50         9.29           AA045608         1.13         7.26         6.45         2.50         9.29           AA628788         4.30         27.63         6.43         1.38         9.13           AA43542         0.26         1.70         6.43         4.03         5.78           AA44864         1.05         6.74         6.41         2.85         6.70           AA75784         0.13         0.21         6.41         5.00         4.97           H15910         0.03         0.21         6.41         5.69         12.25	69	H20847	1.12	7.31	6.53	5.27	9.69	4.63	~
T51617         0.04         0.24         6.51         1.47         11.33           R43328         1.01         6.55         6.50         2.35         11.59           AA953357         1.34         8.71         6.49         10.79         2.65           H21943         1.30         8.44         6.48         4.26         5.16           AA428186         0.02         0.12         6.46         2.50         9.29           AA045608         1.13         7.26         6.45         3.07         8.32           AA62878         4.30         27.63         6.43         1.38         9.13           AA45242         0.07         0.45         6.43         4.03         5.78           AA45542         0.07         0.45         6.41         2.85         6.70           AA45542         0.07         0.45         6.41         2.85         6.70           AA75764         0.13         0.82         6.41         5.00         4.97           H16910         0.03         0.21         6.41         5.69         12.25	733	N48075	0.41	5.64	6.52	4.93	8.97	5.67	7
R43328         1.01         6.55         6.50         2.35         11.69           AA953357         1.34         8.71         6.49         10.79         2.65           H21943         1.30         8.44         6.48         4.26         2.65           AA428186         0.02         0.12         6.46         2.50         9.29           AA628788         1.13         7.26         6.45         3.07         8.32           AA62878         4.30         27.63         6.43         1.38         9.13           AA45242         0.07         0.45         6.43         4.03         5.78           AA452542         0.07         0.45         6.41         2.85         6.70           AA45564         0.13         0.82         6.41         5.00         4.97           H15910         0.03         0.21         6.41         5.69         12.25	<u> </u>	151617	9.0	0.24	6.51	1,47	11.33	6.74	~
AA953357         1.34         8.71         6.49         10.79         2.65           H21943         1.30         8.44         6.48         4.26         5.16           AA428186         0.02         0.12         6.46         2.50         9.29           AA045508         1.13         7.26         6.45         3.07         8.32           AA628788         4.30         27.63         6.43         1.38         9.13           AA034115         0.26         1.70         6.43         4.03         5.78           AA452542         0.07         0.45         6.42         2.05         11.14           AA44864         1.05         8.74         6.41         2.85         6.70           AA455542         0.07         0.45         6.41         5.05         11.14           AA45564         0.13         0.82         6.41         5.09         4.97           H16910         0.03         0.21         6.41         5.69         12.25	8	R43328	1.01	6.55	6.50	2.35	11.59	5.55	7
H21943         1.30         8.44         6.48         4.26         5.16           AA428186         0.02         0.12         6.46         2.50         9.29           AA045508         1.13         7.26         6.45         3.07         8.32           AA628786         4.30         27.63         6.43         1.38         9.13           AA034115         0.26         1.70         6.43         4.03         5.78           AA44864         1.05         6.74         6.41         2.85         6.70           AA757564         0.13         0.82         6.41         5.00         4.97           H16910         0.03         0.21         6.41         5.69         12.25	3251	AA953357	¥.	8.71	6.49	10.79	2.65	9.04	64
AA428186         0.02         0.12         6.46         2.50         9.29           AA045808         1.13         7.26         6.45         3.07         8.32           AA628788         4.30         27.63         6.43         1.38         9.13           AA034115         0.26         1.70         6.43         4.03         5.78           AA45242         0.07         0.45         6.41         2.85         6.70           AA448684         1.05         6.74         6.41         2.85         6.70           AA757584         0.13         0.82         6.41         5.00         4.97           H15910         0.03         0.21         6.41         5.69         12.25	233	H21943	<del>.</del> .	9,44	. 6.48	4.26	5.16	10.01	8
AAG45508         1.13         7.26         6.45         3.07         8.32           AAG28788         4.30         27.63         6.43         1.38         9.13           AAA5242         0.26         1.70         6.43         4.03         5.78           AA45864         1.05         6.74         6.41         2.85         6.70           AA75764         0.13         0.82         6.41         5.00         4.97           H16910         0.03         0.21         6.41         5.69         12.25	264	AA428186	0.02	0.12	6.46	2.50	9.29	7.59	~
AA626788         4.30         27.63         6.43         1.38         9.13           AA034115         0.26         1.70         6.43         4.03         5.78           AA452542         0.07         0.45         6.42         2.05         11.14           AA45664         1.05         6.74         6.41         2.85         6.70           AA757564         0.13         0.82         6.41         5.00         4.97           H15910         0.03         0.21         6.41         5.69         12.25	297	AA045508	1.13	7.26	6.45	3.07	8.32	7.97	8
AA034115 0.26 1.70 6.43 4.03 5.78 AA452542 0.07 0.45 6.42 2.05 11,14 AA448664 1.05 6.74 6.41 2.85 6.70 AA757564 0.13 0.82 6.41 5.00 4.97 H15910 0.03 0.21 6.41 5.69 12.25	784	AA626788	4.30	27.63	6.43	1.38	9.13	8.78	7
AA452542 0.07 0.45 6.42 2.05 11,14 AA448664 1.05 6.74 6.41 2.85 6.70 AA757564 0.13 0.82 6.41 5.00 4.97 H15810 0.03 0.21 6.41 5.69 12.25	200	AA034115	0.26	1.70	6.43	4.03	5.78	9.48	8
AA448664 1.05 6.74 6.41 2.85 6.70 AA757564 0.13 0.82 6.41 5.00 4.97 H15910 0.03 0.21 6.41 5.69 12.25	188	AA452542	0.07	0.45	6.42	2.05	11.14	6.08	~
AA757564 0.13 0.82 6.41 5.00 4.97 H15910 0.03 0.21 6.41 5.69 12.25 13	978	AA448684	56.	6.74	6.41	2.85	6.70	9.70	7
H15810 0.03 0.21 6.41 5.69 12.25	955	AA757564	0.13	0.82	6.41	5.00	4.97	9.25	64
•	462	H15910	63	0.21	6.41	5.69	12.25	1.28	7
					5				

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	7	7	7	7	7	7	7	7	7	7	8	7	7	7	7	7	7	7	7	2	7	7	7	7	7	7	7	2	7	7	7	7	7	
12.22	4.22	8.37	12.04	6.13	11.23	9.86	6.32	8.69	8.17	7.26	8.95	7.88	10.08	13.49	5.30	10.52	5.48	9.06	9.66	6.77	8.79	6.60	7.78	10.29	9.56	9.04	4.83	7.48	7.07	5.03	7.86	7.19	5.37	5.40	
90'9	6.40	7.88	5.53	12.20	6.38	8.03	10.78	6.93	8.05	10.13	2.46	8.82	6.21	5.01	0.27	5.05	12.34	8.92	7.60	0.78	5.59	, 10.69	2.18	7.55	2.69	7.76	5.90	9.73	8.86	£.2	6.77	4.55	8.09	3.03	
0.90	8.52	2.87	1.54	0.76	1.48	1.06	1.79	3.26	7.64	1.45	7.40	2.11	2.51	0.27	13.18	3,19	0.87	0.68	1.35	11.05	4.19	54	8.48	29.0	6.15	1.59	7.62	1.12	2.39	1.52	3.55	6.43	4.66	9.62	
6.39	6.38	6.37	6.37	6.36	6.36	6.31	6.30	6.29	6.29	6.28	6.27	6.27	6.27	6.25	6.25	6.25	6.23	6.22	6.20	6.20	6.19	6.16	6.15	6.13	6.13	6.13	6.12	6.11	6.11	60.9	6.09	6.06	6.04	6.02	4
1.66	4.82	3.76	4.54	1.50	6.46	4.87	51.96	4.53	5.87	2.21	1.37	1.58	0.15	2.29	13.69	2.35	4.11	1.36	12.01	0.08	12.17	0.33	2.57	15.94	12.43	2.89	0,14	2,40	16.02	26.54	4.93	90:0	7.89	96.0	
0.26	0.76	0.59	0.71	0.24	1.01	0.77	8.25	0.72	0.93	0.35	0.22	0.25	0.02	0.37	2.19	0.38	99'0	0.22	1.94	0.01	1.97	9.05	0.42	2.60	2.03	0.47	0.02	0.39	2.62	4.36	0.81	0.01	1.31	0.16	
R89846	AA431753	N39611	AA405533	R93068	AA465223	AA459244	AA042812	N63744	W85878	N63988	N77828	H54659	R44201	N24869	AA430367	R11605	AA701978	R38539	H22568	W31389	AA447662	N56875	AA865924	AA630104	AA430744	AA609891	H18017	R92186	AA430052	AA459106	H96392	AA102068	AA425404	AA620437	
194717	782476	277112	772410	196824	814211	814444	486538	292936	416075	289496	302221	203179	34254	267135	769857	130004	435957	23073	52128	320602	813586	277478	1470195	854701	770992	1032080	50983	195817	781109	814303	256680	489805	773147	951068	
448	449	450	451	452	453	454	455	456	457	458	459	8	461	462	463	464	465	466	467	468	<b>\$</b>	470	471	472	473	474	475	478	477	478	479	480	481	<b>\$</b>	

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 5 fold or more in at least 2 cancer cell lines.

2	7	7	2	~	7	2	7	2	~	~	7	~	~	7	7	~	~	7	7	7	~	7	2	~	7	7	7	7	7	7	7	~	2	7	
5.20	7.25	7.87	7.79	6.52	5.38	6.09	8.76	7.53	2.86	8.22	8.94	7.60	6.64	69.6	8.71	7.58	6.22	5.62	5.26	8.09	6.29	5.11	1.55	8.40	7.45	5.77	10.01	5.36	4.86	1.55	9.5¢	1.87	7.73	7.68	
9.92	9.75	7.62	8.57	9.04	11.46	0.30	7.8	3.15	8.19	8.58	6.81	5.43	2.32	1.00	5.22	5.30	7.27	9.55	10.89	5.86	6.03	10.22	10.45	7.72	7.30	9.02	6.27	9.62	5.66	9.78	6.51	9.35	8.12	5.70	
2.89	0.97	2.44	1.57	2.30	0.97	11.41	1.63	7.03	6.66	0.84	1.87	4.55	8.54	6.78	3.50	4.53	3.92	2.24	1.21	3.39	4.97	1.96	5.27	1.10	2.46	2.42	0.89	2.14	6.56	5.72	1.00	5.83	1.17	3.60	
6.00	5.99	5.98	5.98	5.96	5.93	5.93	. P.	5.90	5.90	5.88	5.88	5.86	5.83	5.82	5.81	5.80	5.80	5.80	5.79	5.78	5.76	5.76	5.76	5.74	5.74	5.73	5.72	5.71	5.69	5.68	5.68	5.68	5.68	5.66	<del>1</del> 5
7.71	8.83	8	2.99	0.55	690	1.85	15.84	3.18	0.36	0.76	2.56	8.00	4.03	90:0	43.39	10.85	32.21	4.83	0.22	13.63	0.16	0.59	4.73	2.62	3.37	6.12	1.35	0.15	4.73	1.58	5.71	2:32	2.35	5.08	
7 79	1.48	9	5 6				2.68	45.0	0.06	0.13	0.43	1.37	0.69	0.01	7.47	1.87	5.55	0.83	90.0	2.36	0.03	0.10	0.82	0.46	0.59	1.07	0.24	0.03	0.83	0.28	1.00	0.51	0.41	0.90	
00458546	AA466363	A 604202	A304202	N38233	AA013463	5110100	N3600B	N29918	N59866	AA778603	N73252	W94009	AA709271	H02158	AA454585	N59136	AA625662	AA465090	AA609914	157359	H01332	AA708605	N62128	R13434	R60752	H10788	H37817	R40481	AA150777	R45952	N72274	N51498	R89363	AA228130	
017710	044080	Pacaca	+07C70	4E0E66	coccnet	200000	. 27.38R	271076	284783	1048949	246549	357531	1343468	150623	809535	287637	745360	814995	1031113	74738	147385	506483	287581	28475	38510	46862	191546	27817	504826	34063	291374	282104	196037	667598	
787	3	}	CO# 907	9 5	ğ ç	8 6	. 60	767	287	167	4	495	496	497	498	499	200	501	502	503	20.5	505	206	205	805	605	510	511	512	513	514	515	516	517	

Table 2-11: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	~	7	7	7	~	7	<b>~</b>	2	7	7	7	7	~	7	7	7	7	7	7	~	<b>7</b>	74	~	~	7	7	7	7	~	~	2	7	7	7	
6.19	5.00	8.55	11.39	2.59	5.48	6.45	7.48	5.99	5.46	2.47	7.67	5.02	8.52	6.45	8.81	6.56	7,85	1.70	7.61	6.30	7.49	7.59	8. 4	5.51	8.69	4.14	7.71	8.77	5.87	727	6.98	8.24	5.55	6.26	
2.36	1.37	7.77	4.0	8.42	4.48	4.88	1.26	10.00	9.88	7.41	2.85	6.93	7.04	8.01	0.09	6.29	7.74	7.64	8.14	8.29	6.97	5.09	6.79	8.15	5.96	5.49	6.69	5.32	5.58	3.79	5.22	6.68	7.08	6.07	
8.40	10.55	0.55	5.02	5.85	6.88	5.49	8.06	0.80	1.43	6.89	6.12	4.68	1.02	2.10	7.57	3.62	0.87	7.11	0.67	1.75	1.86	3.64	1.09	2.62	1.61	6.63	7.84	2.06	4.66	5.05	3.91	1.13	3.39	388	
5.65	5.64	5.62	5.62	5.62	5.62	5.61	5.60	5.60	5.59	5.59	5.55	5.54	5.53	5.52	5.49	5.49	5.49	5.48	5.47	5.45	5.44	5.44	5.44	5.43	5.42	5.42	5.41	5.38	5.37	5.37	5.37	5.35	25.	5.33	16
18.53	6.68	0.25	0.10	90.0	25.20	5,38	17.75	20.0	0.17	8,	3.65	0.40	132.57	0.41	66.94	5.08	11.11	8,93	5.38	0.26	0.05	0.31	1.05	0.45	33.51	5.65	0.59	7.13	1.59	11.60	10.19	16	, c	28.0	;
3.28	1.19	0.05	0.02	0.0	4.49	0.96	3.17	0.0	0.03	0.72	0.66	0.0	23.98	0.07	12.19	0.93	202	.63	0.98	0.05	0.01	90.0	0.19	0.08	6.18	\$	0.11	133	030	2.16	1.80	2 17		. c	?
44699560	AA669218	AA757588	AA427400	W93500	R80790	PCPE0H	AA608730	WI06107	HATATE	TOROT	AAAASSAA	H23983	AABZOAOR	AA907714	AAARTEOR	AA115559	R2230R	Hansas	D00819	AA428514	H96356	AA017213	R89287	AA884428	AA706829	AA59RB03	AA883100	A4458877	AA770383	AA480208	AAGGGGG	AAAAAAA	60630444	AMD/0040	110000
433474	856592	305600	979077	357344	146887	00202	950926	161688	103130	12170A	471707	51465	87878	1505898	Radora	491615	130845	241437	197206	071877	256720	361069	195786	1466998	451805	89R097	1487751	814350	450047	780204	431245	244700	75767	#000 F	24 25
£.	5 £	620	3 5	523		3 6	525	200	3 5	77 2	2 2	2 5	23	3 6	3 E	3 72	3 5	3 4 4 4 4 4 4	24	838	538	25	170	27	543	3	77	3 4	2 2	į	<b>3</b> 5	ξ ξ	2	ָרָה בּינוּ מינוי	}

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

01	~	7	7	8	7	8	7	7	2	7	7	7	7	2	7	7	7	73	7	7	7	8	8	7	7	2	2	8	7	7	~	7	7	7	
6.39	6.03	7.16	7.71	3.90	60 <b>.6</b>	5.20	7.18	5.27	6.95	7.50	7.02	8.43	6.43	7.68	<b>5</b> .64	6.90	5.15	6.97	7.33	7.94	5.93	8.34	5.28	5.63	5.12	7.63	0.32	5.48	5.67	7.92	5.33	6.26	5.79	1.87	
6.23	1.01	7.63	7.20	5.91	5.86	9.61	6.76	8.00	6.68	2.95	2.97	5.07	7.32	5.49	5.99	6.91	9.14	5.27	1.00	0.80	5,12	5.30	8.40	5.03	9.02	5.29	8.82	8.12	8.65	5.51	9.68	3.76	8.28	6.10	•
4.34	8.91	1,15	1.02	6.09	9.0	1.07	1.93	2.58	2.22	5.35	5.79	2.21	1.93	2.51	2.95	1.72	1.23	3.25	7.12	6.70	4.3 4.3	1.68	<b>1</b> .	4.62	1.14	2.35	6.08	1.54	0.82	1.63	3.05	6.00	0.91	7.00	
5.32	5.32	5.31	5.31	5.30	5.30	5.29	5.29	5.29	5.28	5.26	5.26	5.24	5.23	5.23	5.19	5.18	5.17	5.16	5.15	5.15	5.13	5.11	5.10	5.09	5.09	5.09	5.07	5.05	5.04	5.02	5.02	5.01	4.99	4.99	17
35.80	0.05	6.28	2.11	13.64	21.66	4.03	0.23	20.12	0.41	3.68	0.14	0.58	44.95	15.25	0.35	9.47	. 0.51	2.47	0.05	90.0	31.82	0.19	0.43	0.77	32.60	1.98	16.55	40.76	0.22	3.45	23.47	4.55	25.5	8.53	
6.73	100	1.18	0.40	2.57	4.09	0.76	3	3.81	0.08	0.70	80.0	0.1	8.60	2.92	20.0	£.8	0.10	0.48	0.01	0.04	6.20	9.0	90.0	0.15	6.40	0.39	3.26	8.07	9.04	0.69	4.68	0.91	0.91	1.71	
AA620558	B73526	R16676	AA598668	H98822	AA455652	N21334	N57858	T77840	AA628190	H59305	AA879474	AA620814	AA133166	AA169645	AA446748	AA644334	AA426561	AA136060	AA398218	W84774	N66132	AA683338	AA777779	AA885851	AA148505	R71689	AA962541	AA421230	AA490605	N49899	AA504272	AA129552	H61552	AA149637	
951125	166473	1206F4	898258	261714	813997	265102	247089	108395	1055831	204289	1505735	1048685	490753	594517	784126	845602	769146	490060	726658	415806	278483	1293016	448920	1500388	491524	155575	1602209	739247	824109	243653	825411	564803	206052	504279	
553	3	¥ ¥	558	257	. E	558	280	26.	262	563	26.	565	999	267	268	269	570	57.	225	573	574	575	676	21.5	578	679	280	581	582	583	88	585	286	587	

**\$** 

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

8	7	7	7	7	2	2	7	2	7	7	7	7	2	2	7	2	7	7	<b>~</b>	N	7	7	~	8	2	~	7	~	~	7	~	~	~	7
6.61	4.03	5.79	0.97	5.65	5.61	6.75	6.79	2.27	6.33	6.25	5.46	7.16	5.90	6.59	1.00	5.64	5.32	1.00	6.60	7.20	29.62	5.83	6.25	6,47	6.37	8.48	6.21	6.12	5.01	5.80	4.	3.64	5.38	6.39
6.33	5.47	8.07	5.74	8.14	5.83	6.08	6.72	7.27	7.99	6.45	7.62	5.39	5.34	5.47	5.04	5.49	8.39	5.50	5.68	5.26	6.24	6.79	6.82	6.47	5.38	90.0	5.74	5.51	3.86	1.33	5.90	5.23	2.63	5.33
1.99	5.36	0.98	8.10	0.97	3.32	1.91	1.20	5.15	0.37	1.97	1.57	2.09	3.38	2.45	8.44	3.32	0.71	7.91	2,12	1.94	2.51	1.60	1.18	1.30	2.49	99'9	2.27	2.52	5.15	6.88	6.62	5.09	5.94	2.19
4.97	4.96	4.95	4.94	4.92	4.92	4.91	4.90	4.90	4.90	4.89	4.88	4.88	4.87	4.84	4.83	4.81	. 4.	4.80	4.80	4.80	4.79	1.7	4.75	4.75	4.75	4.74	4.74	4.72	4.67	4.67	4.65	4.65	4.65	4.63
1,93	0.75	6.34	0.05	8.75	11.53	0.30	2.11	90.0	8.32	0.71	1.49	34.71	2.66	10.43	9.05	0.90	13.99	0.05	92.90	7.55	24.62	3.4	1.68	09.0	4.36	16.44	0.31	4.54	3.48	0.12	0.27	1.69	1.82	4.18
0.39	0.15	1.28	0.01	1.78	2.34	0.06	0.43	0.02	1.70	0.14	0.30	7.11	5.54	2.16	0.0	0.19	2.91	0.0	19.35	1,57	5.14	0.72	0.35	0.13	0.92	3.47	90.0	0.96	0.74	0.03	90.0	0.36	0.39	0.90
N20480	AA626784	R44346	H39560	AA478542	AA443969	R82041	N56982	AA114106	H16743	AA904969	R55750	AA035384	AA418918	N62372	AA454554	AA156599	AA064627	R43271	AA456931	N31641	AA131526	R54846	H55915	AA521411	R00275	AA779165	AA700415	H05768	166936	AA701587	AA676590	AA046321	H08796	AA443094
264162	197778	34294	186132	784772	757206	.147925	277545	583201	50114	1504152	40721	471598	767994	290560	809508	502296	382451	32517	838568	258966	503715	154472	204129	826089	123255	453005	460553	43826	66406	432732	882488	376764	46097	809466
588	289	280	591	295	593	594	595	296	597	208	599	009	8	602	603	8	605	909	607	809	609	610	611	612	613	614	615	616	617	618	619	620	53	622

Table 2-11; Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line: Genes upregulated 5 fold or more in at least 2 cancer cell lines.

7	7	2	7	6	2	2	7	.~	8	7		8	7	64	6	73	62	63	7	7	2	7	8	~	8	7	7	7	7	7	7	7	2	7
5.88	5.45	6.25	5.98	5.05	7.00	. 228	5.85	5.23	5.23	7.00	5.54	5.34	5.12	5.56	6.95	20.5	5.57	3.12	8.78	1.7	5.93	5.98	5.87	9.76	5.17	5,29	5.41	6,30	5.23	5,54	5.26	5.86	0.44	6.47
5.63	5.08	5.20	5.24	5.72	0.75	2.06	6.62	7.47	7.39	5,53	5.84	5.08	6.30	2.55	5.28	6.17	6.98	5,13	5.09	5.25	6.24	5.02	5.71	5.63	2.84	0.22	5.60	0.31	5.46	5.14	6.04	5.25	6.96	5.51
2.36	3.28	2.32	2.55	2.99	5.99	6.34	1.15	0.93	66'0	1.08	2,21	3.16	2.12	5.38	1.27	2.26	0.90	5.20	1.56	6.45	1.21	2.31	1.72	- 0.87	5.24	7.74	2.19	6.57	2.44	2.38	1.74	26.1	5.57	68:0
4.62	4.60	4.59	4.59	4.59	4.58	4.56	4.54 25.4	4. 2.	4.54	4.54	4.53	4.53	4.52	4.50	4.50	4.49	4.48	4.48	4.48	4.47	4.46	4.	4.44	4.42	4.42	4.42	4.40	4.39	4.38	4.35	4.35	4.34	4.33	4.32
5.17	11.07	15.49	4.45	4.50	0.00	12.03	13.59	5.05	2.67	0.38	2.96	13.59	40.40	2.71	6.39	4.53	1.60	10.51	0.32	1.09	0.53	90'0	0.49	1.88	90.0	7.57	57.53	0.14	0.84	0.73	0.40	0.42	0.10	1.07
1.12	240	3.37	28.0	86	0.0	264	5 6	1	0.59	80.0	0.65	3.00	8.95	0.60	1.42	10.1	0.36	2.35	0.07	0.24	0.12	0.0	0.11	0.42	0.02	1.71	13.07	9.0	0.19	0.17	0.09	0.10	0.02	2,5
A A CORRAT	AA173036	AA598797	AA460634	A A 78 4 7 0 4	AA620254	AA676708	Maggg	AA488979	N51367	AA704858	R24451	AA157787	AA421270	AA885096	H58542	T58434	N29624	AA454947	AA778562	H22854	AA883660	H20743	AA904776	R89765	R06372	AA398355	T90621	AA404338	AA112979	AA884382	AA626247	AA625791	AA443300	AA205320
745107	505400	Section 1	705605	227.745	744044	882355	261567	824728	283208	453229	132954	588840	731031	1468466	206272	75650	257312	814765	1048899	51700	1466599	61328	1504005	194811	126239	726858	110772	758302	530545	1461715	745523	744911	784589	847818
623	624	, S	3 2	22.5	828	20 6	830	8	63	£ 23	75	635	636	637	638	639	9	3	642	543	44	645	98	647	648	649	920	651	652	653	456	855	959	657

ADOVA	170122	5.27	22.74	4.32	1.50	5.60	5.84
757463	AA437245	0.09	0.39	4.28	1.45	5.78	5.62
45514	H08227	90:0	0.27	4.28	1.28	5.77	5.79
280909	N50828	64.1	6.07	4.25	50.1	5.61	6.07
814906	AA465692	2.68	11.37	4.24	1.91	5.44	5.38
272238	N35579	2.10	8.82	4.19	125	5.55	5.76
42400	R60949	1.37	5.73	4.17	5,58	1.07	5.87
281103	N50935	2.75	11.43	4.16	1.40	5.50	5.58
1466824	AA883755	60.0	0.36	4,16	1,38	5.73	5,36
505997	AA708431	0.41	1.69	4.14	5.38	1.63	5.41
841287	AA487206	2.36	9.76	4.13	0.84	6.05	5.49
126695	R07012	0.14	0.57	4.12	1.58	5.14	5.65
858292	AA633993	2.38	9.78	4.10	25.0	5.73	6.01
469211	AA862814	0.20	0.83	4.09	1.23	5.12	5.83
73659	T54527	1.12	4.55	4.08	76.0	5.71	5.55
79726	T62552	<u>5</u> .	4.25	4.08	5.88	5.23	1.12
92388	N79230	1.82	7.30	4.01	5.25	1.78	5.01
239661	H79566	0.02	0.07	3.92	5.37	5.82	0.56
182483	AA676598	2.77	10.81	3.91	1.13	5.04	5.56
49517	H84915	0.76	2.97	3.90	1.12	5.12	5.46
47587	N58136	0.02	0.09	3.90	5.46	5.69	0.55
93420	H47089	0.40	1.56	3.87	1.01	5.60	5.01
87141	AA045340	1.95	2.56	3.87	0.43	5.58	5.63

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostastic Epithelial Cell Line: Genes upregulated 10 fold or more in at least 2 cancer cell lines.

Order	IMAGE	Gen Bank Accession	MPM1(PrEC)	Ave Cancer Coll Lines	Fold (Ave Cell Ine/MPM1)	Fold(MPM2/M F PM1)	Fold(MPN2/M Fold(MPN271/MP Fold(MPN272/M #-of Cancer PM1)	Fold(MPM272/M PW1)	#-of Cancer Cell Lines
	Clone ID	Number		Values	•	37 67	305 13	22.85	n
-	1343732	AA725564	0.01	1.12	112.48	15.43	400	43.45	eo
	78429R	AA447079	0.01	99.0	65.30	43.64	000	75 70	67
•		24400463	50	0.35	63.57	16.10	138.91	23.70	, ,
3	6996/	A4400403	5 6	96 7	54.32	97.06	17.92	47.97	9 (
4	1558655	AA976561	90.0	0.4	ED 47	15.70	123.80	11.93	m
S	743016	AA406036	0.01	0.50	30.4	2 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8	29.19	50.51	က
9	33076	R44048	0.80	36.21	45.45	20.00	70 02	22.90	n
7	626390	AA189106	0.07	3.16	45.16	20.18	6,04	14.51	m
• •	812170	AA456035	0.01	0:30	37.18	16.25	80.73	40.68	m
• •	502AB	H17888	0.01	0.32	36.79	12.25	8/.45	200	· e
, ;	487861	AA045436	0.05	0.59	32.57	29.44	39.44	20.02	, er
2;	2000	N72446	0.23	6.85	29.19	24.48	38.30	24.00	) ("
Ξ !	750187	2720011	190	17.34	27.31	19.07	21.59	41.29	, (
<del>1</del>	263014	H99813	3 8	220	27.21	13.82	36.04	31.77	<b>.</b> .
t.	303180	N92764	200		25.98	33.86	30.16	13.87	W
<del>4</del>	46896	H09818	c0:0	17:1	25.03	38.89	13.26	25.64	ო
5	151184	H02294	0.48	12.53	25.62		52.20	12.48	m
9	41595	R59556	0.01	0.17	25.59	14.00	12 64	28 60	ო
: 5	430082	AA009840	0.40	10.00	24.80	34.18	13:01	40.88	ო
: 9	1049033	44778675	0.38	9.30	24.51	46.38	16.26	20.00	, (r
2 9	005700	8680644	10.0	0.24	24.29	30.26	10.09	32,50	) <b>•</b>
<b>2</b> 2 3	004/80	240230	. 46	3.52	22.24	38.20	16.96	11.55	o •
R 1	365149	AAU23142	0.01	0.25	20.42	18.18	18.45	24.62	, c
5 1	31301	170771	200	0.31	18.24	10.20	24.35	20.16 5.16	, i
73	145/161	AA66510/		2 05	17.83	10.45	28.90	14.13	n 1
ឌ	488431	AA047441	- G	33	15.94	21.48	14.07	12.25	m
75	82869	T69270	0.02	0.32	3 64	13.47	11.86	15.62	m
52	898286	AA598974	1.21	16.52	20.51		R3 14	1917.92	7
<b>5</b> 8	42627	R60995	0.01	9.14	660.46	75.7	GB 37	1701.31	7
72	757165	AA443950	0.01	5.71	591.67	S. 1	90°50	14.43	7
;	592707	AA160606	0.01	1.66	224.30	8.10	680.66	2 T	
3 8	TATOTA	AAAARAAB	0.01	1.70	228.86	10.58	674.66	 	•
3					-				

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

105.59 146.14 105.59 12.54 13.76 10.72 12.17 12.17 12.17 12.17 11.05 11.03 11.01 11.03 10.03 10.	150.46 239.88 229.57 95.72 168.10 4.69 51.92 88.50 101.07	0.89 3.09 1.42 1.42 1.03 1.03 1.03 1.03 1.03 1.03 1.03 1.03	85.17 81.58 65.49 61.20 58.88 56.02 56.02 49.93 44.72 41.11 35.62 34.73 34.73 34.73 36.35 29.15 29.15 27.13 26.99 25.24	0.82 5.16 0.61 12.41 1.10 5.02 13.02 0.44 0.42 0.25 0.33 0.33 0.40 0.40 0.40 0.27 14.38	0.01 0.05 0.05 0.05 0.05 0.05 0.01 0.01	AA418488  W84289  R23270  AA251784  W73144  R44741  R51617  AA504844  R09747  AA504844  R09747  AA608325  H94482  AA608325  AA608322  AA608322  AA608328  AA608328  AA608328  AA608328  AA608328  AA608328  AA608328  AA789328  AA789328  AA789328  AA789328  AA789328  AA789328  AA789338
69.04 27.83 35.03		1.21 0.69 7.36	27.13 26.99 26.24	0.27	0.01 0.03	2 2 2 1
70.25		1.21	27.34	0.27	0.0	~ Z .
5.78	36 8	5.99	28.83	0.44	0.02	m
¥ 4	8 8	9.10 7.73	29.16 29.13	0.40	0.01	
4 % :	8 8	72.0	30.35	7.89	0.26	
2 2	73.	20.19	34.44	0.25	0.01	_
8	62.	0.93	34.73	8.56	0.25	
<b>20</b>	99 90 80 80	4.81 19.5	41.11	0.41	0.01	ю
; <u>8</u>	5 5	4.59	41.72	0.42	0.0	
28	26.	0.91	49.67	13.02	0.26	
8 E	£ 4	0.63 99.52	50.72 49.93	1.10 5.02	0.02 0.10	
22	88.	78.84	56.02	12.41	0.22	
35	51.	1.03	58.88	2.87	0.05	
2 g	4.6	3.73	61.20 59.79	0.61 26.93	0.01	
2	95.	0.87	65.49	5.18	0.08	60
.57	229	1.42	81.58	0.82	0.01	
£ 89	239	3.09	85.17		5.5	
04.46	150	0.89		0.51	č	,
.75	140	0.80	95.78 85.65	1.93 6.83 0.51	0.02	. 4 1

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

N 6	~	7	7	7	7	7	7	7	74	2	7	8	8	2	7	7	8	7	7	7	8	64	2	7	7	7	7	8	7	7	8	7	7	2	)	
11.25	43.13	85.6	48.25	26.43	25.34	31.76	11.14	54.20	23.77	3.08	47.62	22.42	17.48	9.30	28.27	20.17	45.41	40.27	25.28	1.08	43.06	10.71	16.22	10.73	16.88	12.21	34,04	31.84	34.18	33.48	26.82	28.36	25.15	19 22	77:81	
60.48	28.65	17.55	21.91	34.50	42.67	30.37	51.95	10.05	1.50	23.72	10.88	33.42	42.76	30.57	30.55	38.72	10.31	11.18	20.81	14.34	10.12	28.83	34.15	33.82	8 G	37.48	11.78	10.67	15.	90 x	28.4	60.61	10.32	EU.1.2	77.19	
1.76	0.95	44.37	0.87		; ;	3 8	3.82	3 6	8.5	27.50	9 9	ž ų	3 6	5 5	£0.3	67.	3 F	8/7 F	<b>.</b> .	C.33	41.41	6	84.4 70	15.7	30.70	25.83	0. 4	5 6	6.10 47.65	3 5	10,4/	5.81	2.63	0.69	4.69	
24.50	24.24	22 07	18:53	23.00	55.0	23.00	22.37	22.30	Z 15	27.70	\$ 17	20.97	20.40	80.02 50.03	20.27	20.04	19.97	19.50	19.50	19.47	18.93	18,85	18.01	17.44	17.30	17.25	17.19	61.71	16.89	16.78	16.35	16.17	15.84	15.64	15.57	თ
0.24		18.7	0.16 8 :	3.51	0.16	0.23	0.16	3.74	3.65	0.14	40.77	19.16	0.20	6.19	2.46	3.94	0.20	0.20	0.19	3.30	21.01	0.17	0.35	0.17	0.21	33.78	1.06	0.63	0.17	0.1	0.16	11.31	2.94	5.98	0.30	
5	5	0.31	0.01	0.15	0.01	0.0 TO:	0.0	0.17	0.17	0.01	96.1	0.91	0.0	0.30	0.12	0.20	0.01	0.01	0.01	0.17	1.1	0.01	0.02	0.01	0.01	1.96	90:0	0.0 4	0.01	0.01	0.01	0.70	0.19	0.38	0.02	
17000	19004	AA490945	R39191	AA504625	N63057	170198	AA630328	R44936	AA491261	H10072	T61938	N32904	N32542	AA255954	H22563	R95684	AA461508	R86970	R54036	H28091	T65736	R54492	AA057195	AA001376	AA701315	149355	AA857163	N47012	AA676998	AA481481	W92263	W93717	AA504838	AA417713	AA045175	
	251806	824025	23345	825606	284714	81050	855910	34204	824270	46843	79000	259884	270975	686552	51916	199337	795831	197265	40010	162310	80338	39722	489076	361899	435099	67440	1410444	280155	454083	756629	358990	357373	825726	746232	487766	
;	69	99	29	89	69	2	۲	72	2	7	75	76	#	87	78	80	81	82	83	2	. £	8	78	88	28	8	9	8	8	8	92	96	26	: <b>6</b>	8 8	ı

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line, Genes upregulated 10 fold or more in at least 2 cancer cell lines.

7	2	8	7	2	2	2	7	2	7	7	8	7	7	~	7	64	7	7	~	7	2	8	7	7	М	7	7	7	7	7	7	7	7	7
15.13	18.34	15.00	22.27	15.54	23.25	28.45	27.66	4.48	20.74	12.18	17.41	18.27	19.45	18.18	6.86	26.50	25.70	11.26	0.58	26.39	20.98	11.09	17.09	17.55	20.10	12.88	16.24	19.11	16.72	14.01	12.09	3.40	20.99	16.34
28.43	26.10	24.61	16.46	3.02	18.93	4.19	11.04	18.97	9.76	26.44	15.99	21.58	12.04	18.08	17.85	1.54	12.28	21.72	14.55	0.39	14.19	23.87	13.80	15.47	14.95	18.70	17.13	13.18	8.33	20.54	19.31	20.61	10.18	16.85
1.53	0.42	5.14	5.11	24.69	1.00	10.07	2.99	17.85	10.64	2.42	7.33	0.64	8.23	3.39	14.72	10.97	0.65	5.65	22.57	10.40	1.96	2.11	5.65	3.28	1.00	4. 4.	2.08	3.09	10.15	0.58	3.60	10.87	3.65	0.33
15.03	14.95	14.92	14.61	14.41	14.40	14.24	13.89	13.76	13.71	13.68	13.58	13.50	13.24	13.22	13.14	13.00	12.88	12.88	12.56	12.39	12,38	12.36	12.18	12.10	12.02	11.98	11.82	11.80	11.73	11,71	11.67	11.63	11.61	11.17
3.91	3.94	0.35	0.35	1.64	0.14	3.51	4.17	0.13	15.63	1.31	0.17	3.95	8.63	0.26	0.23	90:0	0.13	0.17	0.13	9.44	0.12	3.72	14.39	0.31	1.69	16.82	23.53	3.89	0.31	0.18	11.24	0.20	27.64	0.34
0.26	0.26	0.02	0.02	0.1	0.01	0.25	0.30	0.01	1.14	0.10	0.01	0.29	0.65	0.02	0.02	0.01	0.01	0.01	0.01	92.0	0.01	0.30	1.18	0.03	0.14	1.40	1.99	0.33	0.03	0.02	96.0	0.02	2.38	9.03
AA483445	H18932	H97140	R40434	AA476543	AA434388	AA883800	W90705	R56149	AA628867	R37738	AA128617	AA427719	W83379	AA491227	H19217	N52193	AA017379	N34042	AA453616	AA620697	AA022498	H10713	AA262211	AA778717	R92201	AA706901	AA477227	H09940	AA609002	AA161188	R44816	W93523	AA026682	AA160670
811766	51599	250069	28298	785293	770860	1461609	418004	41406	1035796	26806	502003	771004	415089	824074	51103	284288	381122	244050	795446	1049185	364352	47264	686172	1049079	195845	451648	739450	46461	1030798	591157	33839	357138	366971	592728
100	101	<u>5</u>	ই	፯	105	5	107	801	5	5	111	112	113	<del>1</del>	15	116	117	118	119	120	121	12	52	124	125	126	121	128	129	<del>د</del> ا	131	132	¥	¥

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

<b>,</b> 0	8	7	2	7	7	7	7	7	8	7	7	7	7	7	7	. 2	7	8	~	2	7	7	7	7	2	7	7	23	7	7	7	8	7		
13.80	10.20	12.02	20.10	12.92	7.26	15.60	15.64	1.65	17.42	16.30	14.06	15.31	10.88	7.37	10.89	5.79	11.50	15.44	16.97	15.06	14.44	5.24	4.33	11.30	13.51	13.08	14.35	14.67	5,5	16.60	11.94	14.35	11.27		
17.96	20.27	15.78	10.22	B 10	13.18	14.48	14.23	17.16	11.85	12.15	3.07	13.69	17.62	10.28	15.63	40.79	11.42	1.85	3 5	3 4	12.53	. 27 65	88.0	5.00	3 5	10:01	3 9 0	10.87	13.45	2 0	90.0	56.1	12.48	į	
0.92	0.51	2.03 4.64	4 6 4 5	5.03	11.07	P 80	0.00	2.5	15.0	28.5	42.74	0.67	) ÷	5 5	\$ 50 c	2.63	08.11	9:93	61.4	10.45	2.50	(e,0)	0.00	12.48	4.58	. c. c	04.6	12.50	25.5	\$ 00 of	6.03 5 4	27.11	10.33	<u> </u>	
10.90	10.84	10.83	10.81	10.79	10.70	10.64	10.32	10.21	10.11	10.07	9.85	n d Di d	n t	9.87	9.57	73 (F	9.52	9.48	9.48	9.47	9.33 \$4. 1	9.31	9.24	9.22	9.17	9.12	9.09	5.07	9.06	<b>6</b> .03	8.98	8.87	8.56	8.39	2
3.90	6.12	0.31	0.06	9.32	0.09	13.51	0.87	5.55	0.12	2.53	2.29	1.05	4.94	3.82	0.16	0.77	10,61	1.40	1.40	0.09	90.0	0.22	1,03	76.0	9.89	2.44	0.26	0.26	3.21	11.71	17.81	5.97	0.09	9.51	
0.36	0.57	0.03	0.01	0.86	0.01	1.27	0.08	5.	0.01	0.25	0.23	0.11	0.50	0.39	0.02	0.08	1.12	0.15	0.15	0.01	0.01	0.02	0.11	0.10	1.08	0.27	0.03	0.03	0.35	.3 0	1.98	0.67	0.01	1,13	
H48070	AA410434	N74131	AA461513	N30751	AA453433	. H58234	AA281784	N49389	AA158035	R94504	AA664007	AA625979	AA292226	N21633	AA701652	R46000	N47113_	62406N	AA219060	W16659	AA284072	N50962	H04789	AA705047	AA446462	R96579	R69584	W44766	NZ6515	AA172056	AA620486	W68220	AA496998	W47364	
193333	753428	298417	795840	257955	788209	204483	712401	277621	690640	197651	855391	745490	725877	266135	433567	35804	280375	303088	630013	320495	700792	281162	43733	461488	781047	199577	141726	320865	266259	594758	951241	342640	897546	324717	
135	136	137	138	139	140	141	142	5	₹	145	146	147	148	148	150	151	152	153	45	155	156	157	158	159	160	161	162	<b>1</b>	. <b>⊉</b>	591	<del>8</del>	167	<b>3</b>	69	

Table 2-12: Prostate Cancer Cell Lines vs. Normal Prostatic Epithelial Cell Line. Genes upregulated 10 fold or more in at least 2 cancer cell lines.

7	7	7
12.36	11.26	10.67
11.51	11.14	0.08
1.30	2.23	11.07
8.39	6.21	7.27
3.36	1.49	15.11
0.40	0.18	2.08
AA504265	AA058576	AA706968
825416	380884	451907
170	171	172

9

				•	
Sequence #	Accession #	<b>Dat</b> aba <b>se</b>	Sequence 52	AB033048	GENBANK
Sequence 1	A21185	GENBANK	Sequence 53	AB033114	GENBANK
Sequence 2	AB <b>000509</b>	GENBANK	Sequence 54	AD001528	GENBANK
Sequence 3	AB <b>002329</b>	GENBANK	Sequence 55	AF000986	GENBANK
Sequence 4	AB <b>002357</b>	GENBANK	Sequence 56	AF001893	GENBANK
Sequence 5	AB <b>002366</b>	GENBANK	Sequence 57	AF002715	GENBANK
Sequence 6	AB <b>002387</b>	GENBANK	Sequence 58	AF003837	GENBANK
Sequence 7	AB004304	<b>GEN</b> BANK	Sequence 59	AF004561	GENBANK
Sequence 8	AB <b>004788</b>	GENBANK	Sequence 60	AF006085	GENBANK
Sequence 9	AB <b>004903</b>	GENBANK	Sequence 61	AF007791	GENBANK
Sequence 10	AB <b>007870</b>	GENBANK	Sequence 62	AF008442	GENBANK
Sequence 11	АБ <b>007899</b>	GENBANK .	Sequence 63	AF010235	GENBANK
Sequence 12	AB <b>007930</b>	GENBANK	Sequence 64	AF013988	GENBANK
Sequence 13	AB <b>007941</b>	GENBANK	Sequence 65	AF016266	GENBANK
Sequence 14	AB <b>007956</b>	GENBANK	Sequence 66	AF016270	GENBANK
Sequence 15	AB <b>007957</b>	GENBANK	Sequence 67	AF019226	GENBANK
Sequence 16	AB <b>007960</b>	GENBANK	Sequence 68	AF023259	GENBANK
Sequence 17	AB0 <b>09284</b>	GENBANK	Sequence 69	AF025840	GENBANK
Sequence 18	AB <b>011108</b>	GENBANK	Sequence 70	AF025998	GENBANK
Sequence 19	AB <b>011132</b>	GENBANK	Sequence 71	AF026939	GENBANK
Sequence 20	AB <b>011140</b>	GENBANK	Sequence 72	AF027824	GENBANK
Sequence 21	AB <b>011155</b>	GENBANK	Sequence 73	AF029890	GENBANK
Sequence 22	AB <b>011159</b>	GENBANK	Sequence 74	AF033095	GENBANK
Sequence 23	AB <b>011164</b>	GENBANK	Sequence 75	AF034607	GENBANK
Sequence 24	AB <b>011472</b>	GENBANK	Sequence 76	AF035296	GENBANK
Sequence 25	AB <b>012130</b>	GENBANK	Sequence 77	AF035298	GENBANK
Sequence 26	AB <b>012910</b>	GENBANK	Sequence 78	AF035309	GENBANK
Sequence 27	AB <b>014521</b>	GENBANK	Sequence 79	AF038404	GENBANK
Sequence 28	AB014589	GENBANK	Sequence 80	AF038957	GENBANK
Sequence 29	AB <b>017007</b>	GENBANK	Sequence 81	AF038960	GENBANK
Sequence 30	AB018268	GENBANK	Sequence 82	AF039703	GENBANK
Sequence 31	AB0 <b>18305</b>	GENBANK	Sequence 83	AF039918	GENBANK
Sequence 32	AB018327	GENBANK	Sequence 84	AF042331	GENBANK
Sequence 33	AB018344	GENBANK	Sequence 85	AF044588	GENBANK
Sequence 34	AB018351	GENBANK	Sequence 86	AF045167	GENBANK
Sequence 35	AB019568	GENBANK	Sequence 87	AF045184	GENBANK
Sequence 36	AB020637	GENBANK	Sequence 88	AF047020 AF047472	GENBANK GENBANK
Sequence 37 Sequence 38	AB020669	GENBANK '	Sequence 89	AF048977	GENBANK
Sequence 39	AB0 <b>20680</b> AB0 <b>20682</b>	GENBANK GENBANK	Sequence 90 Sequence 91	AF050171	GENBANK
Sequence 40	AB020692	GENBANK	Sequence 92	AF052124	GENBANK
Sequence 41	AB020092 AB020723	GENBANK	Sequence 92 Sequence 93	AF052164	GENBANK
Sequence 42	AB020723 AB020981	GENBANK	Sequence 94	AF052182	GENBANK
Sequence 43	AB021288	GENBANK	Sequence 95	AF053470	GENBANK
Sequence 44	AB023153	GENBANK	Sequence 96	AF054179	GENBANK
Sequence 45	AB023106	GENBANK	Sequence 97	AF054187	GENBANK
Sequence 45	AB023227	GENBANK	Sequence 98	AF054838	GENBANK
Sequence 47	AB024704	GENBANK	Sequence 99	AF054990	GENBANK
Sequence 48	AB028624	GENBANK	Sequence 100	AF055012	GENBANK
Sequence 49	AB029000	GENBANK	Sequence 101	AF057160	GENBANK
Sequence 50	AB029004	GENBANK	Sequence 101	AF061258	GENBANK
Sequence 51	AB033020	GENBANK	Sequence 102	AF061938	GENBANK
	/ 100000Z0	CEITE/11414	Ocquerioe 105	, 00,000	

Sequence 104	AF067656	GENBANK		Sequence 156	AF155832	GENBANK
Sequence 105	AF067972	GENBANK		Sequence 157	AF159056	GENBANK
Sequence 106	AF068846	GENBANK		Sequence 158	AF176574	GENBANK .
Sequ <b>ence 107</b>	AF069601	GENBANK		Sequence 159	AF182645	GENBANK
Sequence 108	AF070523	GENBANK		Sequence 160	AF188745	GENBANK
Sequence 109	AF070555	GENBANK		Sequence 161	AF192979	GENBANK
Sequence 110	AF070561	GENBANK		Sequence 162	AJ001306	GENBANK
Sequence 111	AF070655	GENBANK		Sequence 163	AJ004913	GENBANK
Sequence 112	AF070657	GENBANK	•	Sequence 164	AJ004955	GENBANK
Sequence 113	AF070664	GENBANK		Sequence 165	AJ006470	GENBANK
Sequence 114	AF071202	GENBANK		Sequence 166	AJ006834	GENBANK
Sequence 115	AF071593	GENBANK		Sequence 167	AJ007398	GENBANK
Sequence 116	AF077030	GENBANK		Sequence 168	AJ010442	GENBANK
Sequence 117	AF077034	GENBANK		Sequence 169	AJ010444	GENBANK
Sequence 118	AF077367	GENBANK		Sequence 170	AJ012499	GENBANK
Sequence 119	AF083441	GENBANK		Sequence 171	AJ130733	GENBANK
Sequence 120	AF083470	GENBANK		Sequence 172	AJ224172	GENBANK
Sequence 121	AF085845	GENBANK		Sequence 173	AJ227918	GENBANK
Sequence 122	AF086003	GENBANK		Sequence 174	AJ228139	GENBANK
Sequence 123	AF086218	GENBANK		Sequence 175	AJ238095	GENBANK
Sequence 124	AF086336	GENBANK		Sequence 176	AJ250042	GENBANK
Sequence 125	AF086431	GENBANK		Sequence 177	AL049227	GENBANK
Sequence 126	AF.086495	GENBANK		Sequence 178	AL049246	GENBANK
Sequence 127	AF086517	GENBANK		Sequence 179	AL049471	GENBANK
Sequence 128	AF086557	GENBANK		Sequence 180	AL049799	GENBANK
Sequence 129	AF087020	GENBANK		Sequence 181	AL049932	GENBANK
Sequence 130	AF087573	GENBANK		Sequence 182	AL049941	GENBANK
Sequence 131	AF088055	GENBANK		Sequence 183	AL049942	GENBANK
Sequence 132	AF091089	GENBANK		Sequence 184	AL049969	GENBANK
Sequence 133	AF093118	GENBANK		Sequence 185	AL049987	GENBANK
Sequence 134	AF100615	GENBANK		Sequence 186	AL050041	GENBANK
Sequence 135	AF100755	GENBANK		Sequence 187	AL050141	GENBANK
Sequence 136	AF100756	GENBANK		Sequence 188	AL050161	GENBANK ·
Sequence 137	AF101051	GENBANK		Sequence 189	AL050184	GENBANK
Sequence 138	AF103774	GENBANK		Sequence 190	AL050198	GENBANK
Sequence 139	AF103907	GENBANK		Sequence 191	AL080172	GENBANK
Sequence 140	AF106681	GENBANK		Sequence 192	AL080234	GENBANK
Sequence 141 Sequence 142	AF110643 AF112152	GENBANK		Sequence 193	AL096719	GENBANK
Sequence 143		GENBANK		Sequence 194	AL110144	GENBANK
Sequence 144	AF114471 AF119297	GENBANK		Sequence 195	AL110183	GENBANK
Sequence 145	AF124438	GENBANK GENBANK		Sequence 196 Sequence 197	AL110197 AL110206	GENBANK
Sequence 146	AF128527	GENBANK		•		GENBANK GENBANK
Sequence 147	AF134159	GENBANK		Sequence 198 Sequence 199	AL110212	GENBANK
Sequence 148	AF139461	GENBANK		•	AL117237	
Sequence 149	AF147380	GENBANK		Sequence 200	AL117427 AL117429	GENBANK GENBANK
Sequence 150	AF151103	GENBANK		Sequence 201	AL117429 AL117505	GENBANK
Sequence 151	AF151103	GENBANK		Sequence 202		
Sequence 151	AF151903	GENBANK		Sequence 203	AL117534	GENBANK
Sequence 153	AF152306	GENBANK		Sequence 204	AL117554 AL117576	GENBANK GENBANK
Sequence 154	AF153191	GENBANK		Sequence 205		GENBANK
Sequence 155	AF155099	GENBANK		Sequence 206	AL117609	
Codamine 100	W. 100099	GLINDAIN		Sequence 207	AL117643	GENBANK

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Sequence 208	AL133000	GENBANK	Sequence 260	J03209	GENBANK
Sequence 209	D00015	GENBANK	•	J03250	GENBANK
Sequence 210	D00017	GENBANK	Sequence 262	J03464	GENBANK
Sequence 211	D13641	GENBANK	Sequence 263	J03779	GENBANK
Sequence 212	D14041	GENBANK	Sequence 264	J03799	GENBANK
Sequence 213	D14696	GENBANK	Sequence 265	J04164	GENBANK
Sequence 214	D16431	GENBANK	Seq <b>uence 266</b>	J04621	GENBANK
Sequence 215	D25215	<b>GEN</b> BANK	Sequ <b>ence 267</b>	J04970	GENBANK
Sequence 216	D26488	<b>GEN</b> BANK	Sequ <b>ence 268</b>	J05021	GENBANK
Sequence 217	D28358	<b>GEN</b> BANK	Sequ <b>ence 269</b>	J05032	· GENBANK
Sequence 218	D28420	<b>GEN</b> BANK	Sequ <b>ence 270</b>	K01911	GENBANK
Sequence 219	D28482	GENBANK	Sequence 271	K02765	GENBANK
Sequence 220	D29954	<b>GEN</b> BANK	Sequ <b>ence 272</b>	L00160	GENBANK
Sequence 221	D29958	GENBANK	Sequence 273	L01439	GENBANK
Sequence 222	D30037	<b>GEN</b> BANK	Sequence 274	L02426	GENBANK
Sequence 223	D31839	GENBANK	Sequ <b>ence 275</b>	L05425	GENBANK
Sequence 224	D37965	GENBANK	Sequ <b>ence 276</b>	L06133	GENBANK
Sequence 225	D38551	GENBANK	Sequence 277	L08441	GENBANK
Sequence 226	D38583	GENBANK	Sequence 278	L10678	GENBANK
Sequence 227	D42040	GENBANK	Sequence 279	L10910	GENBANK
Sequence 228	D43950	<b>GEN</b> BANK	Sequence 280	L11066	GENBANK
Sequence 229	D50371	<b>GEN</b> BANK	Sequence 281	L11932	GENBANK
Sequence 230	D63486	<b>GEN</b> BANK	Sequence 282	L13210	GENBANK
Sequence 231	D63874	GENBANK	Sequence 283	L13385	GENBANK
Sequence 232	D80009	GENBANK	Sequence 284	L13773	GENBANK
Sequence 233	D83032	<b>GEN</b> BANK	Sequence 285	L13799	GENBANK
Sequence 234	D83197	<b>GEN</b> BANK	Sequence 286	L15702	GENBANK
Sequence 235	D83485	<b>GEN</b> BANK	Sequence 287	L19161	GENBANK
Sequence 236	D84557	GENBANK	Sequ <b>ence 288</b>	L19605	GENBANK
Sequence 237	D86228	GENBANK	Sequ <b>ence 289</b>	L19872	GENBANK
Sequence 238	D86322	<b>GEN</b> BANK	Sequ <b>ence 290</b>	L20941	GENBANK
Sequence 239	D87437	<b>GEN</b> BANK	Sequence 291	L22569	GENBANK
Sequence 240	D87444	<b>GEN</b> BANK	Sequ <b>ence 292</b>	L28010	GENBANK
Sequence 241	D87466	<b>GEN</b> BANK	Sequence 293	L33930	GENBANK
Sequence 242	D87666	GENBANK	Sequ <b>ence 294</b>	L36642	GENBANK
Sequence 243	D87667	<b>GEN</b> BANK	Sequ <b>ence 295</b>	L38941	GENBANK
Sequence 244	D87685	<b>GEN</b> BANK	Sequ <b>ence 296</b>	L38961	GENBANK
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Sequence 259	J03040	GENBANK	Sequence 311	M17324	GENBANK
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Sequence 362	U08815	GENBANK	Sequence 413	X05908	GENBANK
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Sequence 52		dbEST	Sequence 579 AAAAAAA
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Sequence 52		dbEST	Seguence For
Sequence 53		dbEST	Sequence Egg A A Second
Sequence 53		dbEST	Sequence 582 AA244400
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· Sequence 53		dbEST	Seguence ESE AACABOR
Sequence 535		dbEST	Sequence 596 AAGOS (SE
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Sequence 537		dbEST	Sequence FDD
Sequence 538		dbEST	Soguenes Eng
Sequence 539		dbEST	Conversion
Sequence 540		dbEST	Socilores For
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Page 18

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Page 19

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Page 21

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Page 22

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Page 23

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Page 24

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Page 25

### Table 3-1

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Page 27

Sequence#	Accession#	Database	s	equence 2857	AB019524	GENBANK
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Page 28

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Sequence 2924	AF030310	GENBANK	Sequence 2976	AF081282	GENBANK
Sequence 2925	AF033095	GENBANK	Sequence 2977	AF082868	GENBANK
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Sequence 2927	AF038440	GENBANK	Sequence 2979		GENBANK
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Sequence 2930	AF041259	GENBANK	Sequence 2982	AF085893	GENBANK
Sequence 2931	AF042166	GENBANK	Sequence 2983	AF085917	GENBANK
Sequence 2932	AF042385	GENBANK	Sequence 2984	AF086183	GENBANK
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Sequence 2959	AF069765	GENBANK	Sequence 3011	AF145029	GENBANK
Sequence 2960	AF070523	GENBANK	Sequence 3012	AF145316	GENBANK
			Ocquence 30 12	-1-2210	GENDAIN

Page 29

#### Table 3-1

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Sequence 3031	AF176702	GENBANK	Sequence 3083	D16111	GENBANK
Sequence 3032	AF179274	GENBANK	Sequence 3084	D16937	GENBANK
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Page 31

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Page 32

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Page 33

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Page 34

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Page 37

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Page 38

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Page 39

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Page 40

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Sequence 4201	A1017959	dbEST	Sequence 4253	Al141189	dbEST
Sequence 4202	AI018617	dbEST	Sequence 4254	Al141583	dbEST
Sequence 4203	AI024907	dbEST	Sequence 4255	Al141732	<b>dbEST</b>
Sequence 4204	AI025376	dbEST	Sequence 4256	Al148251	dbEST
Sequence 4205	Al027516	dbEST	Sequence 4257	Al150619	dbEST
Sequence 4206	A1034024	dbEST	Sequence 4258	Al160426	dbEST
Sequence 4207	AI038048	dbEST	Sequence 4259	Al161253	dbEST
Sequence 4208	AI039227	dbEST	Sequence 4260	AI167801	dbEST

Page 41

Sequence 4261	Al184320	dbEST	Sequence 4313	Al267677	dbEST
Sequence 4262	AJ189339	dbEST	Sequence 4314	Al267808	dbEST
Sequence 4263	Al189345	dbEST	Sequence 4315	Al267941	dbEST
Sequence 4264	Al189386	dbEST	Sequence 4316	Al269883	dbEST
Sequence 4265	Al189894	dbEST	Sequence 4317	AJ271740	dbEST
Sequence 4266	Al190648	dbEST	Sequence 4318	Al274786	dbEST
Sequence 4267	Al192780	dbEST	Sequence 4319	Al275114	dbEST
Sequence 4268	Al192784	dbEST	Sequence 4320	Al275175	dbEST
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Sequence 4270	Al198311	dbEST	Sequence 4322	Al277739	dbEST
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Sequence 4273	Al199641	dbEST	Sequence 4325	Al284011	dbEST
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Sequence 4275	Al199773	dbEST	Sequence 4327	A1288388	dbEST
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Sequence 4277	Al200444	dbEST	Sequence 4329	Al288929	dbEST
Sequence 4278	Al201261	dbEST	Sequence 4330	Al289120	dbEST
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Sequence 4280	Al203350	dbEST	Sequence 4332	Al292299	dbEST
Sequence 4281	AI207650	dbEST	Sequence 4333	Al306464	dbEST
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Sequence 4299	Al253367	dbEST	Sequence 4351	AI356718	dbEST
Sequence 4300	Al253436	dbEST	Sequence 4352	AI360B70	dbEST
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Sequence 4302	Al262139	dbEST	Sequence 4354	Al362490	dbEST
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Sequence 4305	Al267185	dbEST	Sequence 4357	Al366376	dbEST
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Sequence 4309	AI267502	dbEST	Sequence 4361	Al381996	dbEST
Sequence 4310	Al267574	dbEST	Sequence 4362	Al394173	dbEST
Sequence 4311	Al267596	dbEST	Sequence 4363	Al394498	dbEST
Sequence 4312	AI267652	dbEST	Sequence 4364	Al417833	
			00400100 7004	<del>/ (17</del> 17033	dbEST

Page 42

Sequence 4365	· · · · · · · · · · · · · · · · · · ·	dbEST	Sequence 4417	Al684039	dbEST
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Sequence 4367		dbEST	Sequence 4419	AI686957	dbEST
Sequence 4368	AI434639	dbEST	Sequence 4420	AI690236	dbEST
Sequence 4369	Al436714	dbEST	Sequence 4421	Al692940	dbEST
Sequence 4370	AJ439173	dbEST	Sequence 4422	Al692995	dbEST
Sequence 4371	AJ453358	dbEST	Sequence 4423	AI694572	dbEST
Sequence 4372	Al473779	dbEST	Sequence 4424	AI697815	dbEST
Sequence 4373	AJ476254	dbEST	Sequence 4425	AJ701119	dbEST
Sequence 4374	Al499378	dbEST	Sequence 4426	AI708995	dbEST
Sequence 4375	Al499439	dbEST	Sequence 4427	Al732541	dbEST
Sequence 4376	Al499933	dbEST	Sequence 4428	Al732557	dbEST
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Sequence 4378	AI525426	dbEST	Sequence 4430	AI745280	
Sequence 4379	AI525773	dbEST	Sequence 4431	AI751583	dbEST
Sequence 4380	AI525843	dbEST	Sequence 4432	Al751914	dbEST
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Sequence 4383	AI547284	dbEST	Sequence 4435	Al754013	dbEST
Sequence 4384	AI547309	dbEST	Sequence 4436	A1755085	dbEST
Sequence 4385	AI557226	dbEST	Sequence 4437	Al761508	dbEST
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Sequence 4396	AI589633	dbEST	Sequence 4448	Al815829	dbEST
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Sequence 4415	Al679629	dbEST	Sequence 4466	AI887517	dbEST
Sequence 4416	AI683183	dbEST	Sequence 4467	AI887664	dbEST
		40501	Sequence 4468	AI911703	dbEST

Page 43

### Table 3-1

Sequence 4469	Al914133	dbEST	Sequence 4521	AW026680	dbEST
Sequence 4470	Al917649	dbEST	Sequence 4522	AW027007	dbEST
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Sequence 4473	AI929453	dbEST	Sequence 4525	AW058559	dbEST
Sequence 4474	Al934153	dbEST	Sequence 4526	AW062376	dbEST
Sequence 4475	Al934595	dbEST	Sequence 4527	AW071906	dbEST
Sequence 4476	Al935291	dbEST	Sequence 4528	AW075433	dbEST
Sequence 4477	AI935816	dbEST	Sequence 4529	AW084950	dbEST
Sequence 4478	A1937060	dbEST	Sequence 4530	AW129510	dbEST
Sequence 4479	Al951021	dbEST	Sequence 4531	AW132009	dbEST
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Sequence 4481	A1954800	dbEST	Sequence 4533	AW134867	dbEST
Sequence 4482	Al962736	dbEST	Sequence 4534	AW151254	dbEST
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Sequence 4484	Al973218	dbEST	Sequence 4536	AW160669	dbEST
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Sequence 4488	AL035985	dbEST	Sequence 4540	AW163441	dbEST
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Sequence 4491	AL036445	dbEST	Sequence 4543	AW170485	dbEST
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Sequence 4519	AW016611	dbEST	Sequence 4571	H06040	dbEST
Sequence 4520	AW026543	dbEST	Sequence 4572	H06313	dbEST
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Sequence 457		dbEST	Sequence 4625	R70116	dbEST
Sequence 457/ Sequence 457/	4 H10932	dbEST	Sequence 4626		
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Sequence 4576	H12254	dbEST	Sequence 4628		dbEST
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Sequence 4578		dbEST	Sequence 4630		dbEST
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Sequence 4624	R59031	dbEST	Sequence 4675		NUCPATENT
4024	R61601	dbEST	Sequence 4676	V69855	NUCPATENT
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Page 45

Converse ACTT	\ <i>G</i> 0254	AUTODATENT	S		
Sequence 4677	V70354	NUCPATENT	Sequence 4727	AD00000	GENDANA
Sequence 4678	V72211	NUCPATENT	Sequence 4727	AB002353	GENBANK
Sequence 4679	V81414	NUCPATENT	Sequence 4728	AB002375	GENBANK
Sequence 4680	V84510	NUCPATENT	Sequence 4729	AB002380	GENBANK
Sequence 4681	V86232	NUCPATENT	Sequence 4730	AB002386	GENBANK
Sequence 4682	V89565	NUCPATENT	Sequence 4731	AB007929	GENBANK
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Page 47

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Sequence 4887	M15885	GENBANK	Sequence 4939	U82256	GENBANK
Sequence 4888	M16247	GENBANK	Sequence 4940	U94586	GENBANK
Sequence 4889	M17885	GENBANK	Sequence 4941	X01037	GENBANK
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Sequence 4892	M22538	. GENBANK	Sequence 4944	X04106	GENBANK
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Sequence 4894	M26325	GENBANK	Sequence 4946	X04526	<b>GENBANK</b>
Sequence 4895	M26663	GENBANK	Sequence 4947	X13839	GENBANK
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Sequence 4898	M33308	GENBANK	Sequence 4950	X58141	GENBANK
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Sequence 4904	M64241	GENBANK	Sequence 4956	X69711	GENBANK
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Sequence 4909	S50015	GENBANK	Sequence 4961	X84908	GENBANK
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Page 48 . (of 73 pages in Table 2.1)

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Sequence 4988	AA172239	dbEST	Sequence 5040	AA344067	dbEST
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Sequence 5004	AA256401	dbEST	Sequence 5056	AA443463	dbEST
Sequence 5005	AA258916	dbEST	Sequence 5057	AA446877	dbEST
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Sequence 5037	AA339065	dbEST	Sequence 5089	AA578265	dbEST
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Page 49

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Page 50

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Page 51

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Sequence 5314	AF034209	GENBANK	Sequence 5366	AF078854	GENBANK
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Sequence 5319	AF038451	GENBANK	Sequence 5371	AF083190	GENBANK
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Page 52

Sequence 5401         AF151861         GENBANK         Sequence 5452         D29840         GENBANK           Sequence 5401         AF151884         GENBANK         Sequence 5453         D2984         GENBANK           Sequence 5402         AF151883         GENBANK         Sequence 5454         D31886         GENBANK           Sequence 5403         AF151893         GENBANK         Sequence 5455         D38037         GENBANK           Sequence 5404         AF159056         GENBANK         Sequence 5456         D38549         GENBANK           Sequence 5406         AF159092         GENBANK         Sequence 5457         D42039         GENBANK           Sequence 5407         AF168956         GENBANK         Sequence 5469         D42047         GENBANK           Sequence 5409         AF177777         GENBANK         Sequence 5460         D42138         GENBANK           Sequence 5411         AF188746         GENBANK         Sequence 5460         D43972         GENBANK           Sequence 5411         AF188746         GENBANK         Sequence 5462         D49372         GENBANK           Sequence 5411         AF188746         GENBANK         Sequence 5463         D49489         GENBANK           Sequence 5412         A1				•		
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Sequence 5407         AF159295         GENBANK         Sequence 5459         D42047         GENBANI           Sequence 5408         AF177791         GENBANK         Sequence 5460         D42138         GENBANI           Sequence 5409         AF177775         GENBANK         Sequence 5461         D43950         GENBANI           Sequence 5411         AF188298         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5411         AF188748         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5413         AF180167         GENBANK         Sequence 5464         D50369         GENBANI           Sequence 5413         AF190167         GENBANK         Sequence 5465         D50371         GENBANI           Sequence 5414         AJ010482         GENBANK         Sequence 5465         D50371         GENBANI           Sequence 5415         AJ012499         GENBANK         Sequence 5468         D50919         GENBANI           Sequence 5417         AJ012499         GENBANK         Sequence 5468         D50919         GENBANI           Sequence 5418         AJ130733         GENBANIK         Sequence 5470         D63486         GENBANI           Sequence 5418		AF159056	GENBANK			
Sequence 5407         AF168956         GENBANK         Sequence 5460         D42047         GENBANK           Sequence 5409         AF1777291         GENBANK         Sequence 5460         D42138         GENBANI           Sequence 5409         AF1777275         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5411         AF188298         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5412         AF188746         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5412         AF188746         GENBANK         Sequence 5463         D49489         GENBANI           Sequence 5413         AF190167         GENBANK         Sequence 5465         D50371         GENBANI           Sequence 5413         AJ1010482         GENBANK         Sequence 5466         D50579         GENBANI           Sequence 5415         AJ101482         GENBANK         Sequence 5466         D50579         GENBANI           Sequence 5417         AJ012499         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5417         AJ120733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5420		AF159092	GENBANK			
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Sequence 5409         AF177775         GENBANK         Sequence 5461         D43950         GENBANI           Sequence 5410         AF188298         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5411         AF188746         GENBANK         Sequence 5462         D49489         GENBANI           Sequence 5412         AF188746         GENBANK         Sequence 5462         D50369         GENBANI           Sequence 5413         AF190167         GENBANK         Sequence 5465         D50371         GENBANI           Sequence 5414         AJ010046         GENBANK         Sequence 5466         D50579         GENBANI           Sequence 5416         AJ010482         GENBANK         Sequence 5466         D50919         GENBANI           Sequence 5416         AJ012463         GENBANK         Sequence 5468         D50919         GENBANI           Sequence 5418         AJ1012499         GENBANIK         Sequence 5468         D50919         GENBANI           Sequence 5418         AJ130733         GENBANIK         Sequence 5469         D55653         GENBANI           Sequence 5419         AJ223075         GENBANIK         Sequence 5471         D64110         GENBANI           Sequence 5429         <	Sequence 5407	AF168956	GENBANK	•		
Sequence 5410         AF188298         GENBANK         Sequence 5462         D49372         GENBANI           Sequence 5411         AF188746         GENBANK         Sequence 5463         D49489         GENBANI           Sequence 5412         AF188748         GENBANK         Sequence 5464         D50369         GENBANI           Sequence 5413         AF190167         GENBANK         Sequence 5466         D50371         GENBANI           Sequence 5414         AJ010482         GENBANK         Sequence 5467         D50916         GENBAN           Sequence 5416         AJ012463         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5417         AJ012463         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5417         AJ012493         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5419         AJ223350         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5471         D80099         GENBAN           Sequence 5421         AJ249731	Sequence 5408	AF177291	GENBANK	•		GENBANK
Sequence 5411         AF188745         GENBANK         Sequence 5463         D49489         GENBANK           Sequence 5412         AF188746         GENBANK         Sequence 5464         D50369         GENBANK           Sequence 5413         AF190167         GENBANK         Sequence 5465         D50371         GENBANK           Sequence 5414         AJ010466         GENBANK         Sequence 5466         D50579         GENBAN           Sequence 5416         AJ012463         GENBANK         Sequence 5467         D50916         GENBAN           Sequence 5417         AJ012499         GENBANK         Sequence 5469         D55653         GENBAN           Sequence 5419         AJ233075         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5421         AJ223350         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5421         AJ2249731         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5422         AL049969         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5425         AL05018<		AF177775	GENBANK	•		7
Sequence 5412         AF188746         GENBANK         Sequence 5464         D50369         GENBANK           Sequence 5413         AF190167         GENBANK         Sequence 5465         D50371         GENBANK           Sequence 5414         AJ010046         GENBANK         Sequence 5467         D50579         GENBANK           Sequence 5415         AJ010482         GENBANK         Sequence 5467         D50579         GENBANK           Sequence 5416         AJ012463         GENBANK         Sequence 5467         D50316         GENBAN           Sequence 5417         AJ012493         GENBANK         Sequence 5469         D55653         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5419         AJ223375         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5421         AJ223172         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ2247731         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5423         AL04970	Sequence 5410	AF188298	GENBANK	·		GENBANK
Sequence 5413         AF190167         GENBANK         Sequence 5465         D50371         GENBANK           Sequence 5414         AJ010046         GENBANK         Sequence 5466         D50579         GENBANK           Sequence 5415         AJ010482         GENBANK         Sequence 5467         D50916         GENBAN           Sequence 5416         AJ012463         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5417         AJ012499         GENBANK         Sequence 5469         D55653         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5419         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5421         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5422         AL049736         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5422         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050188 </td <td>Sequence 5411</td> <td>AF188745</td> <td>GENBANK</td> <td>•</td> <td></td> <td>GENBANK</td>	Sequence 5411	AF188745	GENBANK	•		GENBANK
Sequence 5414         AJ010046         GENBANK         Sequence 5466         D50579         GENBANK           Sequence 5415         AJ010482         GENBANK         Sequence 5467         D50916         GENBANK           Sequence 5416         AJ012463         GENBANK         Sequence 5468         D50919         GENBANK           Sequence 5417         AJ012499         GENBANK         Sequence 5469         D55653         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5420         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224172         GENBANK         Sequence 5472         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL050198         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198<	Sequence 5412	AF188746	GENBANK	Sequence 5464		GENBANK
Sequence 5415         AJ010482         GENBANK         Sequence 5467         D50916         GENBAN           Sequence 5416         AJ012463         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5417         AJ012499         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5421         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224731         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5422         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5426         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL080089 <td>Sequence 5413</td> <td>AF190167</td> <td>GENBANK</td> <td>•</td> <td></td> <td>GENBANK</td>	Sequence 5413	AF190167	GENBANK	•		GENBANK
Sequence 5416         AJ012463         GENBANK         Sequence 5468         D50919         GENBAN           Sequence 5417         AJ012499         GENBANK         Sequence 5469         D55653         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5420         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224172         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224172         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL049906         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5475         D85181         GENBAN           Sequence 5426         AL050274 <td>Sequence 5414</td> <td>AJ010046</td> <td></td> <td>Sequence 5466</td> <td>D50579</td> <td>GENBANK</td>	Sequence 5414	AJ010046		Sequence 5466	D50579	GENBANK
Sequence 5417         AJ012499         GENBANK         Sequence 5469         D55653         GENBAN           Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5419         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224721         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135 <td>Sequence 5415</td> <td>AJ010482</td> <td>GENBANK</td> <td>Sequence 5467</td> <td>D50916</td> <td>GENBANK</td>	Sequence 5415	AJ010482	GENBANK	Sequence 5467	D50916	GENBANK
Sequence 5418         AJ130733         GENBANK         Sequence 5470         D63486         GENBAN           Sequence 5419         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224172         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL050274         GENBANK         Sequence 5479         D87127         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87453         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080089 <td>Sequence 5416</td> <td>AJ012463</td> <td>GENBANK</td> <td>•</td> <td>D50919</td> <td>GENBANK</td>	Sequence 5416	AJ012463	GENBANK	•	D50919	GENBANK
Sequence 5419         AJ223075         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224172         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5476         D83735         GENBAN           Sequence 5424         AL049705         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5423         AL050198         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5426         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL080089         GENBANK         Sequence 5479         D87127         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5480         D87452         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87462         GENBAN           Sequence 5430         AL080880 <td>Sequence 5417</td> <td>AJ012499</td> <td>GENBANK</td> <td>Sequence 5469</td> <td>D55653</td> <td>GENBANK</td>	Sequence 5417	AJ012499	GENBANK	Sequence 5469	D55653	GENBANK
Sequence 5419         AJ223375         GENBANK         Sequence 5471         D64110         GENBAN           Sequence 5420         AJ223350         GENBANK         Sequence 5472         D80009         GENBAN           Sequence 5421         AJ224172         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5480         D87666         GENBAN           Sequence 5430         AL110867 <td>Sequence 5418</td> <td>AJ130733</td> <td>GENBANK</td> <td>• •</td> <td></td> <td>GENBANK</td>	Sequence 5418	AJ130733	GENBANK	• •		GENBANK
Sequence 5421         AJ224172         GENBANK         Sequence 5473         D82060         GENBAN           Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBAN           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBAN           Sequence 5426         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 54479         D87453         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBAN           Sequence 5430         AL096880         GENBANK         Sequence 5481         D87666         GENBAN           Sequence 5431         AL109672 <td></td> <td>AJ223075</td> <td>GENBANK</td> <td>Sequence 5471</td> <td>D64110</td> <td>GENBANK</td>		AJ223075	GENBANK	Sequence 5471	D64110	GENBANK
Sequence 5422         AJ249731         GENBANK         Sequence 5474         D83032         GENBANK           Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBANK           Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBAN           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBAN           Sequence 5431         AL109672         GENBANK         Sequence 5482         D87684         GENBAN           Sequence 5432         AL110183         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5433         AL117413 </td <td>Sequence 5420</td> <td>AJ223350</td> <td>GENBANK</td> <td>•</td> <td>D80009</td> <td>GENBANK</td>	Sequence 5420	AJ223350	GENBANK	•	D80009	GENBANK
Sequence 5423         AL049705         GENBANK         Sequence 5475         D83735         GENBAN           Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBAN           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBAN           Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87452         GENBAN           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBAN           Sequence 5431         AL109672         GENBANK         Sequence 5483         D87684         GENBAN           Sequence 5432         AL110183         GENBANK         Sequence 5484         D87735         GENBAN           Sequence 5433         AL117413         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5434         AL117550 <td>Sequence 5421</td> <td>AJ224172</td> <td>GENBANK</td> <td>Sequence 5473</td> <td></td> <td>GENBANK</td>	Sequence 5421	AJ224172	GENBANK	Sequence 5473		GENBANK
Sequence 5424         AL049969         GENBANK         Sequence 5476         D85181         GENBANK           Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBANK           Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBANK           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBAN           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBAN           Sequence 5431         AL109672         GENBANK         Sequence 5483         D87684         GENBAN           Sequence 5432         AL110183         GENBANK         Sequence 5484         D87735         GENBAN           Sequence 5433         AL117413         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5434         AL117550         GENBANK         Sequence 5485         D88674         GENBAN           Sequence 5436         AL121733<	Sequence 5422	AJ249731		Sequence 5474	D83032	GENBANK
Sequence 5425         AL050198         GENBANK         Sequence 5477         D86984         GENBANK           Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBANK           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBANK           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBANK           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBANK           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBANK           Sequence 5431         AL109672         GENBANK         Sequence 5483         D87684         GENBANK           Sequence 5432         AL110183         GENBANK         Sequence 5484         D87735         GENBAN           Sequence 5433         AL117413         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5434         AL117550         GENBANK         Sequence 5486         D88674         GENBAN           Sequence 5435         AL117733         GENBANK         Sequence 5487         D90373         GENBAN           Sequence 5437         D0042	Sequence 5423	AL049705	GENBANK	Sequence 5475	D83735	GENBANK
Sequence 5426         AL050274         GENBANK         Sequence 5478         D87127         GENBAN           Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBAN           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBAN           Sequence 5431         AL109672         GENBANK         Sequence 5483         D87684         GENBAN           Sequence 5432         AL110183         GENBANK         Sequence 5484         D87735         GENBAN           Sequence 5433         AL117413         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5434         AL117550         GENBANK         Sequence 5486         D88674         GENBAN           Sequence 5435         AL117595         GENBANK         Sequence 5487         D90373         GENBAN           Sequence 5436         AL121733         GENBANK         Sequence 5489         D0427         GENBAN           Sequence 5438         D13641	Sequence 5424	AL049969	GENBANK	Sequence 5476	D85181	GENBANK
Sequence 5427         AL080089         GENBANK         Sequence 5479         D87292         GENBAN           Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBAN           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBAN           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBAN           Sequence 5431         AL109672         GENBANK         Sequence 5483         D87684         GENBAN           Sequence 5432         AL110183         GENBANK         Sequence 5484         D87735         GENBAN           Sequence 5433         AL117413         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5434         AL117550         GENBANK         Sequence 5486         D88674         GENBAN           Sequence 5435         AL117595         GENBANK         Sequence 5487         D90373         GENBAN           Sequence 5436         AL121733         GENBANK         Sequence 5488         D90427         GENBAN           Sequence 5437         D00422         GENBANK         Sequence 5489         E01650         GENBAN           Sequence 5438         D13641	Sequence 5425	AL050198	GENBANK	Sequence 5477	D86984	GENBANK
Sequence 5428         AL080135         GENBANK         Sequence 5480         D87453         GENBANK           Sequence 5429         AL080209         GENBANK         Sequence 5481         D87462         GENBANK           Sequence 5430         AL096880         GENBANK         Sequence 5482         D87666         GENBANK           Sequence 5431         AL109672         GENBANK         Sequence 5483         D87684         GENBANK           Sequence 5432         AL110183         GENBANK         Sequence 5484         D87735         GENBAN           Sequence 5433         AL117413         GENBANK         Sequence 5485         D87969         GENBAN           Sequence 5434         AL117550         GENBANK         Sequence 5486         D88674         GENBAN           Sequence 5435         AL117595         GENBANK         Sequence 5487         D90373         GENBAN           Sequence 5436         AL121733         GENBANK         Sequence 5487         D90373         GENBAN           Sequence 5437         D00422         GENBANK         Sequence 5489         E01650         GENBAN           Sequence 5438         D13641         GENBANK         Sequence 5490         E01915         GENBAN           Sequence 5443         D13900	Sequence 5426	AL050274	GENBANK	Sequence 5478	D87127	GENBANK
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Sequence 5442 D14664 GENBANK Sequence 5494 J00194 GENBAN Sequence 5443 D16111 GENBANK Sequence 5495 J02642 GENBAN	Sequence 5440	D13900	GENBANK			
Sequence 5443 D16111 GENBANK Sequence 5495 J02642 GENBAN		D14662	GENBANK	•		GENBANK
Oddonos o 10 11 Oznaka wa	Sequence 5442	D14664	GENBANK			GENBANK
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Codesino Citio Citio Carretta	Sequence 5445	D17409		•		GENBANK
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Page 54

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Page 55

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Page 56

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Page 57

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Page 58

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Page 59

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Page 60

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Page 61

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Page 62

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Sequence 6486	AI080267	dbEST		AI203141	dbEST
Sequence 6487	A1080485	dbEST	Sequence 6538	Al204308	dbEST
Sequence 6488	Al081291	dbEST	Sequence 6539	Al205228	dbEST
Sequence 6489	AI084101	dbEST	Sequence 6540	Al206063	dbEST
Sequence 6490	Al084916	dbEST	Sequence 6541	Al207781	dbEST
7		GDE 31	Sequence 6542	Al216969	dbEST

Page 63

				·		
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Se	quence 6544	Al217172	dbEST	Sequence 6596	Al349099	dbEST
Sec	quence 6545	Al217964	dbEST	Sequence 6597	Al349937	dbEST
Sec	quence 6546	Al219278	dbEST	Sequence 6598	Al357472	dbEST
Se	quence 6547	Al220051	dbEST	Sequence 6599	AJ360165	dbEST
Se	quence 6548	Al239506	dbEST	Sequence 6600	Al366549	dbEST
Se	quence 6549	Al241117	dbEST	Sequence 6601	Al368743	dbEST
Sec	quence 6550	A1241659	dbEST	Sequence 6602	AI369849	dbEST
Sec	quence 6551	Al241744	dbEST	Sequence 6603	Al377012	dbEST
Sec	quence 6552	Al246383	dbEST	Sequence 6604	Al378613	dbEST
Sec	quence 6553	Al249764	· dbEST	Sequence 6605	AJ380537	dbEST
Sec	quence 6554	Al250055	dbEST	Sequence 6606	AI380842	dbEST
Sec	quence 6555	Al252084	dbEST	Sequence 6607	Al401771	dbEST
Sec	quence 65 <b>56</b>	Al252283	dbEST	Sequence 6608	Al417016	dbEST
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Sec	quence 6558	Al253330	dbEST	Sequence 6610	Al431350	dbEST
Sec	quence 6559	Al253335	dbEST	Sequence 6611	Al433818	dbEST
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	uence 6581	Al318091	dbEST	Sequence 6633	Al557246	dbEST
	uence 6582	Al335028	dbEST	Sequence 6634	A1557269	dbEST
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	•	Al336800	dbEST	Sequence 6637	Al563921	dbEST
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	uence 6592	Al342937	dbEST	Sequence 6644	AI630985	dbEST
	uence 6593	Al343692	dbEST	Sequence 6645	AI631745	dbEST
seq	uence 6594	Al346657	dbEST	Sequence 6646	A1632244	dbEST

Page 64

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Sequence 6698	Al917110	dbEST	Sequence 6750	AW160696	dbEST
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Page 65

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Sequence 6792	N39674	dbEST	Sequence 6844	W73290	dbEST
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Sequence 6801	N63387	dbEST	Sequence 6853	T74203	NUCPATENT
Sequence 6802	N75468	dbEST	Sequence 6854	V07906	NUCPATENT

Page 66

Sequence 6855	V30916	NUCPATENT	Sequence 6905	AF035839	GENBANK
Sequence 6856	V34207	NUCPATENT	Sequence 6906	AF044957	GENBANK
Sequence 6857	V34226	NUCPATENT	Sequence 6907	AF048977	GENBANK
Sequence 6858	V34305	NUCPATENT	Sequence 6908	AF052138	GENBANK
Sequence 6859	V35555	NUCPATENT	Sequence 6909	AF054183	GENBANK
Sequence 6860	V43604	NUCPATENT	Sequence 6910	AF054987	GENBANK
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Sequence 6862	V58756	NUCPATENT	Sequence 6912	AF055474	GEN <b>BANK</b>
Sequence 6863	V58761	NUCPATENT	Sequence 6913	AF067396	GENBANK
Sequence 6864	V62427	NUCPATENT	Sequence 6914	AF068117	GENBANK
Sequence 6865	V69855	NUCPATENT	Sequence 6915	AF068754	GENBANK
Sequence 6866	V70354	NUCPATENT	Sequence 6916	AF069301	GENBANK
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Sequence 6868	V72211	NUCPATENT	Sequence 6918	AF071172	GENBANK
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Sequence 6891	AC33416	PREPATNUC	Sequence 6941	AL050291	GENBANK
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Sequence 6902	AB015644	GENBANK	Sequence 6953	D87666	GENBANK
Sequence 6903	AB016087	GENBANK	Sequence 6954	D88153	GENBANK
Sequence 6904	AF020351	GENBANK	Sequence 6955	E01915	GENBANK

Page 67

Sequence 6956	J02642	GENBANK	Sequence 7007	AA042813	dbEST
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Sequence 6958	J04208	GENBANK	Sequence 7009	AA045753	dbEST
Sequence 6959	L035 <b>58</b>	GENBANK	Sequence 7010	AA045879	dbEST
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Sequence 6961	L14778	GENBANK	Sequence 7012	AA062804	dbEST
Sequence 6962	L20422	GENBANK	Sequence 7013	AA062928 -	dbEST
Sequence 6963	M11147	GENBANK	Sequence 7014	AA064627	dbEST
Sequence 6964	M11560	GENBANK	Sequence 7015	AA070911	dbEST
Sequence 6965	M17885	GENBANK	Sequence 7016	AA076077	dbEST
Sequence 6966	M37583	GENBANK	Sequence 7017	AA076593	dbEST
Sequence 6967	M38188	GENBANK	Sequence 7018	AA081815	dbEST
Sequence 6968	M64241	GENBANK	Sequence 7019	AA082502	dbEST
Sequence 6969	U12979	GENBANK	Sequence 7020	AA083481	dbEST
Sequence 6970	U17104	GENBANK	Sequence 7021	AA083794	dbEST
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Page 68

# Table 3-1

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Sequence 7063		dbEST	Sequence 7114	AA393336	dbEST
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Sequence 7065		dbEST	Sequence 7116		dbEST
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Sequence 7068		dbEST	Sequence 7119	AA401237	dbEST
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Sequence 7070	AA203462	dbEST	Sequence 7121	AA404535	dbEST
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Sequence 7210	AA604815	dbEST	Sequence 7261	AA827816	dbEST

Page 70

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Page 71

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Page 72

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Sequence 7508	X24921	NUCPATENT
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Sequence 7511	X89652	NUCPATENT
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Page 73

Sequence 1240

Sequence 2802

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Sequence 2803

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Sequence 2804

Sequence 2805

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Sequence 4710

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Sequence 4711

Sequence 4712

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#### Sequence 4713

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TTGGAATGGTGGGCACAGTGGCTGTCACTGTCATGCCTCAGTGGAGAGTGTCGGCCTTCA
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GGCAGGCTAACATCAGGATGCAGTTGCAAAATCTATGATTCCCTGCTGGCTCTTTCTCCG
GACCTACAGGCAGCCAGAGGACTGATGTTGCGCTGCTTCCGTGATGTCCTTTCTCTCCG

#### Sequence 4714

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#### Sequence 4715

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#### Sequence 4716

#### Sequence 4717

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#### Sequence 4718

Sequence 4719

Sequence 4720

Sequence 4722

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Sequence 6893

Sequence 6894

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ACCCTCTGGTAATGAGGCGGGGAAGAAGGAGGGGGAGGCGAGGGGTGACATCGTGGGAGAC
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Sequence 6895

Sequence 6896

Sequence #	Acc No	Database hit	GI Number
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Sequence 7513	A26481	Genbank	1247452
Sequence 7514	A26595	Genbank	1247460
Sequence 7515	A29216	Genbank	1247528
Sequence 7516	A33292	Genbank	1567852
Sequence 7517	A36460	Genbank	2293778
Sequence 7518	A43521	Genbank	2298710
Sequence 7519	A58331	Genbank	3713993
Sequence 7520	A80239	Genbank	6092971
Sequence 7521	A81324	Genbank	6731650
Sequence 7522	A98295	Genbank	6781400
Sequence 7523	AB000360	Genbank	2547041
Sequence 7524	AB000468	Genbank	1843400
Sequence 7525	AB000516	Genbank	2723379
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Sequence 7528	AB000889	Genbank	2467299
Sequence 7529	AB001523	Genbank	2244605
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Sequence 7533	AB002299	Genbank	2224542
Sequence 7534	AB002310	Genbank	2224564
Sequence 7535	AB002313	Genbank	2280475
Sequence 7536	AB002323	Genbank	2224590
Sequence 7537	AB002328	Genbank	2224600
Sequence 7538	AB002336	Genbank	2224616
Sequence 7539	AB002346	Genbank	6634018
Sequence 7540	AB002351	Genbank	2224646
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Sequence 7547	AB002377	Genbank	6634024
Sequence 7548 Sequence 7549	AB002381	Genbank	2224706
Sequence 7550	AB002382	Genbank	2224708
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Sequence 7552	AB002559 AB003476	Genbank Genbank	1944129
Sequence 7553	AB003730	Genbank	2081606
Sequence 7554	AB005754	Genbank	2627128 6456482
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Sequence 7563	AB007900	Genbank	2662160
Sequence 7564	AB007920	Genbank	3413863
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Sequence 7566	AB007931	Genbank	3413885
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Sequence 7567         AB007957         Genbank         3413931           Sequence 7569         AB007965         Genbank         3413931           Sequence 7570         AB009398         Genbank         3618342           Sequence 7571         AB011086         Genbank         3043551           Sequence 7573         AB011096         Genbank         3043551           Sequence 7574         AB0111086         Genbank         3043551           Sequence 7575         AB0111149         Genbank         3043667           Sequence 7576         AB0111151         Genbank         3043667           Sequence 7578         AB0111151         Genbank         3043681           Sequence 7578         AB0111151         Genbank         3043681           Sequence 7578         AB0111166         Genbank         3043713           Sequence 7580         AB011167         Genbank         3043713           Sequence 7581         AB011167         Genbank         3043713           Sequence 7583         AB011536         Genbank         3043717           Sequence 7584         AB011542         Genbank         3043729           Sequence 7585         AB0121293         Genbank         3449297				
Sequence 7569         AB007965         Genbank         3413940           Sequence 7571         AB0019398         Genbank         3618342           Sequence 7571         AB011087         Genbank         3043551           Sequence 7573         AB011096         Genbank         3043551           Sequence 7574         AB011113         Genbank         3043667           Sequence 7576         AB011144         Genbank         3043667           Sequence 7578         AB011151         Genbank         3043681           Sequence 7578         AB011155         Genbank         3043681           Sequence 7578         AB011166         Genbank         3043681           Sequence 7579         AB011167         Genbank         3043713           Sequence 7581         AB011167         Genbank         3043713           Sequence 7582         AB011167         Genbank         3043712           Sequence 7583         AB011156         Genbank         3043713           Sequence 7584         AB0111542         Genbank         3043729           Sequence 7585         AB012124         Genbank         3449297           Sequence 7586         AB012124         Genbank         342608           Sequen	Sequence 756	7 AB007944	Genbank	3413911
Sequence 7569         AB007965         Genbank         3413940           Sequence 7571         AB009398         Genbank         3018342           Sequence 7572         AB011087         Genbank         3043553           Sequence 7573         AB011096         Genbank         3043551           Sequence 7574         AB011113         Genbank         3043667           Sequence 7576         AB011144         Genbank         3043667           Sequence 7578         AB011155         Genbank         3043681           Sequence 7578         AB011155         Genbank         3043681           Sequence 7578         AB011166         Genbank         3043713           Sequence 7579         AB011167         Genbank         3043713           Sequence 7581         AB011167         Genbank         3043713           Sequence 7582         AB011167         Genbank         3043713           Sequence 7583         AB011156         Genbank         3043713           Sequence 7584         AB0111542         Genbank         3043729           Sequence 7585         AB012124         Genbank         3449297           Sequence 7586         AB012124         Genbank         342609           Sequenc	Sequence 756	B AB007957	Genbank	3413931
Sequence 7570         AB009398         Genbank         3043551           Sequence 7571         AB011086         Genbank         3043551           Sequence 7573         AB011096         Genbank         3043553           Sequence 7574         AB0111113         Genbank         3043605           Sequence 7576         AB0111149         Genbank         3043607           Sequence 7577         AB011151         Genbank         3043681           Sequence 7578         AB011151         Genbank         3043688           Sequence 7579         AB011165         Genbank         3043713           Sequence 7580         AB011167         Genbank         3043713           Sequence 7581         AB011169         Genbank         3043713           Sequence 7582         AB011175         Genbank         3043713           Sequence 7584         AB011536         Genbank         3043729           Sequence 7584         AB011542         Genbank         3449297           Sequence 7586         AB012124         Genbank         3449293           Sequence 7586         AB012124         Genbank         3449309           Sequence 7586         AB012129         Genbank         3362605           Seque				
Sequence 7571         AB011086         Genbank         3043551           Sequence 7572         AB011087         Genbank         3043551           Sequence 7573         AB011096         Genbank         3043571           Sequence 7574         AB011113         Genbank         3043605           Sequence 7576         AB011144         Genbank         3043667           Sequence 7577         AB011155         Genbank         3043689           Sequence 7578         AB011165         Genbank         3043681           Sequence 7579         AB011166         Genbank         3043711           Sequence 7581         AB011167         Genbank         3043711           Sequence 7581         AB011156         Genbank         3043717           Sequence 7581         AB011156         Genbank         3043717           Sequence 7583         AB011536         Genbank         3043729           Sequence 7584         AB011542         Genbank         3449297           Sequence 7586         AB012124         Genbank         3449290           Sequence 7586         AB012913         Genbank         349930           Sequence 7586         AB012913         Genbank         3527083           Sequence			Genbank	
Sequence 7572         AB011087         Genbank         3043553           Sequence 7573         AB011096         Genbank         3043671           Sequence 7576         AB011114         Genbank         3043667           Sequence 7576         AB011149         Genbank         3043667           Sequence 7577         AB011151         Genbank         3043689           Sequence 7578         AB011166         Genbank         3043711           Sequence 7580         AB011167         Genbank         3043713           Sequence 7581         AB011169         Genbank         3043713           Sequence 7582         AB011176         Genbank         3043713           Sequence 7583         AB011536         Genbank         3043713           Sequence 7584         AB011536         Genbank         3043729           Sequence 7583         AB011542         Genbank         3449297           Sequence 7586         AB012124         Genbank         3449297           Sequence 7586         AB012129         Genbank         349399           Sequence 7586         AB01291         Genbank         302802           Sequence 7588         AB01492         Genbank         3327083           Sequence 75				
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Sequence 7575         AB011144         Genbank         3043667           Sequence 7576         AB0111151         Genbank         3043677           Sequence 7577         AB011151         Genbank         3043681           Sequence 7579         AB011166         Genbank         3043713           Sequence 7580         AB011167         Genbank         3043713           Sequence 7581         AB011169         Genbank         3043713           Sequence 7582         AB011175         Genbank         3043713           Sequence 7583         AB011536         Genbank         3043729           Sequence 7584         AB011536         Genbank         3449297           Sequence 7585         AB012124         Genbank         3449309           Sequence 7586         AB012124         Genbank         3599675           Sequence 7587         AB01291         Genbank         3062802           Sequence 7588         AB012921         Genbank         3062802           Sequence 7589         AB014087         Genbank         307093           Sequence 7589         AB014087         Genbank         3327083           Sequence 7591         AB014535         Genbank         3327127           Sequence				
Sequence 7576         AB011149         Genbank         3043677           Sequence 7577         AB011151         Genbank         3043681           Sequence 7578         AB011166         Genbank         3043711           Sequence 7580         AB011167         Genbank         3043713           Sequence 7581         AB011169         Genbank         3043713           Sequence 7582         AB011175         Genbank         3043729           Sequence 7584         AB011536         Genbank         3043729           Sequence 7584         AB011542         Genbank         3449297           Sequence 7586         AB012124         Genbank         4126408           Sequence 7586         AB012912         Genbank         362802           Sequence 7588         AB012911         Genbank         362802           Sequence 7588         AB01487         Genbank         362802           Sequence 7589         AB014535         Genbank         3327083           Sequence 7590         AB014540         Genbank         3327093           Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014566         Genbank         3327145           Sequence 75				
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Sequence 7578         AB011155         Genbank         3043689           Sequence 7579         AB011166         Genbank         3043711           Sequence 7580         AB011167         Genbank         3043717           Sequence 7581         AB011175         Genbank         3043729           Sequence 7583         AB011536         Genbank         3449297           Sequence 7584         AB011542         Genbank         3449309           Sequence 7585         AB012124         Genbank         3599675           Sequence 7587         AB012913         Genbank         3599675           Sequence 7587         AB012911         Genbank         3062802           Sequence 7587         AB012911         Genbank         3062802           Sequence 7588         AB012922         Genbank         3062802           Sequence 7589         AB014087         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327083           Sequence 7592         AB014542         Genbank         3327193           Sequence 7593         AB014567         Genbank         3327127           Sequence 7594         AB014569         Genbank         3327145           Sequenc				
Sequence 7579         AB011166         Genbank         3043711           Sequence 7580         AB011167         Genbank         3043713           Sequence 7581         AB011169         Genbank         3043729           Sequence 7583         AB011536         Genbank         3043729           Sequence 7584         AB011542         Genbank         3449297           Sequence 7585         AB012124         Genbank         4126408           Sequence 7586         AB012912         Genbank         3599675           Sequence 7586         AB012911         Genbank         362802           Sequence 7588         AB012911         Genbank         362802           Sequence 7588         AB012922         Genbank         362802           Sequence 7590         AB01487         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327093           Sequence 7593         AB014542         Genbank         3327197           Sequence 7594         AB014563         Genbank         3327127           Sequence 7595         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327191           Sequence 75				
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Sequence 7581         AB011169         Genbank         3043729           Sequence 7583         AB011175         Genbank         3043729           Sequence 7584         AB011536         Genbank         3449297           Sequence 7585         AB011542         Genbank         3449309           Sequence 7586         AB012124         Genbank         3199675           Sequence 7587         AB012191         Genbank         3062802           Sequence 7588         AB012911         Genbank         3062802           Sequence 7588         AB012921         Genbank         5672599           Sequence 7589         AB014087         Genbank         5672599           Sequence 7591         AB014540         Genbank         3327093           Sequence 7593         AB014542         Genbank         3327097           Sequence 7593         AB014563         Genbank         3327127           Sequence 7594         AB014566         Genbank         3327127           Sequence 7595         AB014569         Genbank         3327191           Sequence 7597         AB014589         Genbank         3327191           Sequence 7598         AB014597         Genbank         3327207           Sequenc				
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Sequence 7583         AB011536         Genbank         3449297           Sequence 7584         AB011542         Genbank         3449309           Sequence 7586         AB012124         Genbank         3599675           Sequence 7587         AB012911         Genbank         3062802           Sequence 7588         AB012922         Genbank         4126426           Sequence 7589         AB014087         Genbank         5672599           Sequence 7590         AB014535         Genbank         3327093           Sequence 7591         AB014540         Genbank         3327093           Sequence 7592         AB014540         Genbank         3327093           Sequence 7593         AB014557         Genbank         3327127           Sequence 7593         AB014563         Genbank         3327127           Sequence 7595         AB014569         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327191           Sequence 7597         AB014589         Genbank         3327197           Sequence 7600         AB015597         Genbank         3327197           Sequence 7601         AB017314         Genbank         3582440           Sequenc				
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Sequence 7585         AB012124         Genbank         3599675           Sequence 7587         AB012911         Genbank         3599675           Sequence 7588         AB012911         Genbank         3062802           Sequence 7589         AB014087         Genbank         5672599           Sequence 7590         AB014535         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327093           Sequence 7592         AB014542         Genbank         3327097           Sequence 7593         AB014563         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327145           Sequence 7595         AB014569         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327191           Sequence 7597         AB014589         Genbank         3327191           Sequence 7598         AB014597         Genbank         3327197           Sequence 7600         AB015597         Genbank         4586879           Sequence 7601         AB017498         Genbank         3582440           Sequence 7603         AB017548         Genbank         3885244           Sequenc				
Sequence 7586         AB012193         Genbank         3599675           Sequence 7587         AB012911         Genbank         3062802           Sequence 7588         AB012922         Genbank         4126426           Sequence 7599         AB014087         Genbank         5672599           Sequence 7590         AB014535         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327097           Sequence 7592         AB014542         Genbank         3327097           Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327191           Sequence 7598         AB014592         Genbank         3327191           Sequence 7599         AB014597         Genbank         3327197           Sequence 7601         AB017314         Genbank         4536879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB018247         Genbank         3885244           Sequenc				
Sequence 7587         AB012911         Genbank         3062802           Sequence 7588         AB012922         Genbank         4126426           Sequence 7589         AB014087         Genbank         5672599           Sequence 7590         AB014535         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327093           Sequence 7592         AB014542         Genbank         3327127           Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014566         Genbank         3327145           Sequence 7595         AB014569         Genbank         3327151           Sequence 7596         AB014589         Genbank         3327197           Sequence 7598         AB014592         Genbank         3327197           Sequence 7599         AB014597         Genbank         3327197           Sequence 7600         AB015597         Genbank         4586879           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582144           Sequence 7604         AB018247         Genbank         3885363           Sequenc				
Sequence 7588         AB012922         Genbank         5672599           Sequence 7590         AB014087         Genbank         5672599           Sequence 7591         AB014535         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327093           Sequence 7592         AB014542         Genbank         3327127           Sequence 7593         AB014563         Genbank         3327127           Sequence 7594         AB014566         Genbank         3327145           Sequence 7595         AB014569         Genbank         3327151           Sequence 7596         AB014589         Genbank         3327191           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014592         Genbank         3327197           Sequence 7600         AB015597         Genbank         3327207           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         4586879           Sequence 7603         AB017498         Genbank         3882144           Sequence 7604         AB018247         Genbank         3882144           Sequenc			Genbank	3599675
Sequence 7589         AB014087         Genbank         3327083           Sequence 7591         AB014535         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327097           Sequence 7592         AB014542         Genbank         3327097           Sequence 7593         AB014563         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327145           Sequence 7595         AB014569         Genbank         3327151           Sequence 7596         AB014589         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014597         Genbank         3327197           Sequence 7600         AB015597         Genbank         3327197           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3882144           Sequence 7605         AB018247         Genbank         3882144           Sequence 7606         AB018255         Genbank         3882144           Sequenc			Genbank	3062802
Sequence 7590         AB014535         Genbank         3327083           Sequence 7591         AB014540         Genbank         3327097           Sequence 7592         AB014542         Genbank         3327097           Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014569         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327191           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014592         Genbank         3327207           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         415713           Sequence 7601         AB017144         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3882144           Sequence 7604         AB018247         Genbank         3882144           Sequence 7606         AB018255         Genbank         3882144           Sequence	Sequence 7588	B AB012922	Genbank	4126426
Sequence 7591         AB014540         Genbank         3327093           Sequence 7592         AB014542         Genbank         3327097           Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014569         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327151           Sequence 7697         AB014589         Genbank         3327191           Sequence 7598         AB014592         Genbank         3327197           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB018247         Genbank         3882144           Sequence 7605         AB018247         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882214           Sequenc	Sequence 7589	9 AB014087	Genbank	5672599
Sequence 7592         AB014542         Genbank         3327097           Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014566         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         3327207           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882214           Sequence 7607         AB018302         Genbank         3882231           Sequenc	Sequence 759	O AB014535	Genbank	3327083
Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014566         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014592         Genbank         3327207           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7609         AB018302         Genbank         3882214           Sequence 7610         AB018315         Genbank         3882238           Sequenc	Sequence 759	1 AB014540	Genbank	3327093
Sequence 7593         AB014557         Genbank         3327127           Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014566         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014592         Genbank         3327207           Sequence 7600         AB015597         Genbank         3327207           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582144           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         3882144           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequenc	Sequence 7592	2 AB014542	Genbank	3327097
Sequence 7594         AB014563         Genbank         3327139           Sequence 7595         AB014566         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327197           Sequence 7598         AB014592         Genbank         3327207           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882144           Sequence 7609         AB018302         Genbank         3882214           Sequence 7610         AB018315         Genbank         3882238           Sequenc			Genbank	3327127
Sequence 7595         AB014566         Genbank         3327145           Sequence 7596         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327191           Sequence 7598         AB014592         Genbank         3327197           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882144           Sequence 7609         AB018302         Genbank         3882214           Sequence 7610         AB018315         Genbank         3882238           Sequence 7611         AB018316         Genbank         3882266           Sequenc			Genbank	3327139
Sequence 7596         AB014569         Genbank         3327151           Sequence 7597         AB014589         Genbank         3327191           Sequence 7598         AB014592         Genbank         3327197           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7609         AB018302         Genbank         3882214           Sequence 7610         AB018315         Genbank         3882238           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018331         Genbank         3882282           Sequenc			Genbank	
Sequence 7597         AB014589         Genbank         3327191           Sequence 7598         AB014592         Genbank         3327197           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         3882144           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018331         Genbank         3882282           Sequenc				
Sequence 7598         AB014592         Genbank         3327197           Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882266           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018331         Genbank         3882266           Sequence 7614         AB018331         Genbank         3882304           Sequenc				
Sequence 7599         AB014597         Genbank         3327207           Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882266           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882266           Sequence 7613         AB018331         Genbank         3882304           Sequence 7614         AB018335         Genbank         3882318           Sequenc				
Sequence 7600         AB015597         Genbank         4115713           Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018315         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882304           Sequence 7614         AB018335         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882334           Sequenc				
Sequence 7601         AB017114         Genbank         4586879           Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018315         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882304           Sequence 7614         AB018335         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequenc				
Sequence 7602         AB017335         Genbank         3582440           Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018315         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882304           Sequence 7614         AB018335         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018357         Genbank         3882340           Sequenc				
Sequence 7603         AB017498         Genbank         3582144           Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018315         Genbank         3882266           Sequence 7612         AB018316         Genbank         3882262           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018357         Genbank         3882340           Sequenc				
Sequence 7604         AB017548         Genbank         3885363           Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018315         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018357         Genbank         3882340           Sequence 7620         AB019564         Genbank         3885367           Sequenc				
Sequence 7605         AB018247         Genbank         4049592           Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018353         Genbank         3882340           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7606         AB018255         Genbank         3882144           Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018353         Genbank         3882340           Sequence 7620         AB018357         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7607         AB018281         Genbank         3882196           Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7608         AB018290         Genbank         3882214           Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882340           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7609         AB018302         Genbank         3882238           Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7610         AB018315         Genbank         3882264           Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7611         AB018316         Genbank         3882266           Sequence 7612         AB018324         Genbank         3882282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7612         AB018324         Genbank         3862282           Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7613         AB018331         Genbank         3882296           Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7614         AB018335         Genbank         3882304           Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7615         AB018342         Genbank         3882318           Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371			·	
Sequence 7616         AB018347         Genbank         3882328           Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7617         AB018350         Genbank         3882334           Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7618         AB018353         Genbank         3882340           Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7619         AB018357         Genbank         6009486           Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7620         AB019564         Genbank         3885367           Sequence 7621         AB019568         Genbank         3885371				
Sequence 7621 AB019568 Genbank 3885371				
_ · · ·				
Sequence 7622 AB020315 Genbank 3925828				
	Sequence 7622	2 AB020315	Genbank	3925828

Table 3-3

Sequence 7623	AB020335	Genbank	6518494
Sequence 7624	AB020623	Genbank	3985929
Sequence 7625	AB020631	Genbank	4240136
Sequence 7626	AB020645	Genbank	4240164
Sequence 7627	AB020659	Genbank	4240192
Sequence 7628	AB020662	Genbank	4240198
Sequence 7629	AB020666	Genbank	6635134
Sequence 7630	AB020674	Genbank	4240222
Sequence 7631	AB020679	Genbank	4240232
Sequence 7632	AB020692	Genbank	4240258
Sequence 7633	AB020697	Genbank	4240268
Sequence 7634	AB020698	Genbank	4240270
Sequence 7635	AB020713	Genbank	4240270
Sequence 7636	AB020713	Genbank	4240300
Sequence 7637	AB020714 AB020718	Genbank	
Sequence 7638	AB020718 AB020721	Genbank	4240310
Sequence 7639	AB020721 AB020764		4240316
		Genbank	6467144
Sequence 7640	AB020874	Genbank	4003394
Sequence 7641	AB021663	Genbank	4996450
Sequence 7642	AB022537	Genbank	6942217
Sequence 7643	AB023049	Genbank	5672604
Sequence 7644	AB023050	Genbank	5672605
Sequence 7645	AB023051	Genbank	5672606
Sequence 7646	AB023052	Genbank	5672607
Sequence 7647	AB023137	Genbank	4589471
Sequence 7648	AB023145	Genbank	4589487
Sequence 7649	AB023150	Genbank	4589509
Sequence 7650	AB023152	Genbank	<b>458951</b> 3
Sequence 7651	AB023160	Genbank	4589529
Sequence 7652	AB023165	Genbank	4589539
Sequence 7653	AB023195	Genbank	4589599
Sequence 7654	AB023199	Genbank	4589607
Sequence 7655	AB023205	Genbank	4589619
Sequence 7656	AB023420	Genbank	4579908
Sequence 7657	AB024704	Genbank	4589928
Sequence 7658	AB025580	Genbank	<b>59315</b> 68
Sequence 7659	AB026906	Genbank	5295993
Sequence 7660	AB028893	Genbank	6552364
Sequence 7661	AB028942	Genbank	<b>56893</b> 68
Sequence 7662	AB028952	Genbank	5689394
Sequence 7663	AB028985	Genbank	5689460
Sequence 7664	AB028986	Genbank	5689462
Sequence 7665	AB028990	Genbank	5689470
Sequence 7666	AB028999	Genbank	5689488
Sequence 7667	AB029002	Genbank	5689494
Sequence 7668	AB029003	Genbank	5689496
Sequence 7669	AB029008	Genbank	5689506
Sequence 7670	AB029013	Genbank	5689516
Sequence 7671	AB029017	Genbank	5689524
Sequence 7672	AB029019	Genbank	5689528
Sequence 7673	AB029025	Genbank	5689540
Sequence 7674	AB029026	Genbank	5689542
Sequence 7675	AB029027	Genbank	5689544
Sequence 7676	AB031051	Genbank	6683742
Sequence 7677	AB032417	Genbank	6277265
Sequence 7678	AB032963	Genbank	6329896

Sequence 7679	AB032975	Genbank	6330044
Sequence 7680	AB032978	Genbank	6330094
Sequence 7681	AB032983	Genbank	6330128
Sequence 7682	AB033007	Genbank	6330242
Sequence 7683	AB033011	Genbank	6330301
		Genbank	6330350
Sequence 7684	AB033018		
Sequence 7685	AB033023	Genbank	6330384
Sequence 7686	AB033026	Genbank	6330406
Sequence 7687	AB033027	Genbank	6330415
Sequence 7688	AB033033	Genbank	6330496
Seguence 7689	AB033051	Genbank	6330667
Sequence 7690	AB033058	Genbank	6330721
Sequence 7691	AB033070	Genbank	6330818
Sequence 7692	AB033073	Genbank	6330839
Sequence 7693	AB033078	Genbank	6330873
Sequence 7694	AB033104	Genbank	6331314
Sequence 7695	AB034205	Genbank	6899845
	AB035519	Genbank	6681348
Sequence 7696			
Sequence 7697	AB035698	Genbank	6970477
Sequence 7698	AB037728	Genbank	7242968
Sequence 7699	AB037743	Genbank	7243024
Sequence 7700	AB037748	Genbank	7243034
Sequence 7701	AB037761	Genbank	7243060
Sequence 7702	AB037763	Genbank	7243064
Sequence 7703	AB037775	Genbank	7243088
Sequence 7704	AB037783	Genbank	7243104
Sequence 7705	AB037784	Genbank	7243106
Sequence 7706	AB037797	Genbank	7243132
Sequence 7707	AB037801	Genbank	7243140
Sequence 7708	AB037807	Genbank	7243152
Sequence 7709	AB037819	Genbank	7243176
	AB037844	Genbank	7243226
Sequence 7710		Genbank	7243220
Sequence 7711	AB037847		
Sequence 7712	AB037856	Genbank	7243267
Sequence 7713	AC000024	Genbank	2995601
Sequence 7714	AC000031	Genbank	4646246
Sequence 7715	AC000036	Genbank	7958979
Sequence 7716	AC000049	Genbank	5705995
Sequence 7717	AC000083	Genbank	3810636
Sequence 7718	AC000087	Genbank	7124755
Sequence 7719	AC000088	Genbank	7124834
Sequence 7720	AC000095	Genbank	7940365
Sequence 7721	AC000114	Genbank	1809233
Sequence 7722	AC000134	Genbank	4755212
Sequence 7723	AC000353	Genbank	6970735
Sequence 7724	AC000378	Genbank	2270906
Sequence 7725	AC002064	Genbank	2076723
Sequence 7726	AC002066	Genbank	2076721
		Genbank	
Sequence 7727	AC002094		2155224
Sequence 7728	AC002105	Genbank	2288973
Sequence 7729	AC002117	Genbank	2281075
Sequence 7730	AC002347	Genbank	2828783
Sequence 7731	AC002377	Genbank	2275192
Sequence 7732	AC002401	Genbank	2599240
Sequence 7733	AC002470	Genbank	6478944
Sequence 7734	AC002522	Genbank	7124014

Table 3-3

Sequence 7735	AC002550	Genbank	2570261
Sequence 7736	AC002553	Genbank	3126783
Sequence 7737	AC003010	Genbank	2981254
Sequence 7738	AC003025	Genbank	3337308
Sequence 7739	AC003031	Genbank	2570227
Sequence 7740	AC003036	Genbank	2935595
Sequence 7741	AC003098	Genbank	2822155
Sequence 7742	AC003103	Genbank	2842782
Sequence 7743	AC003663	Genbank	3097871
Sequence 7744	AC003665	Genbank	3808091
Sequence 7745	AC003687	Genbank	2828778
Sequence 7746	AC003950	Genbank	4262275
Sequence 7747	AC003963	Genbank	2995604
Sequence 7748	AC004016	Genbank	2781374
Sequence 7749	AC004017	Genbank	2795792
Sequence 7750	AC004020	Genbank	3219330
Sequence 7751	AC004080	Genbank	2822164
Sequence 7752	AC004084	Genbank	2822156
Sequence 7753	AC004098	Genbank	3097872
Sequence 7754	AC004126	Genbank	3242744
Sequence 7755	AC004150	Genbank	4585948
Sequence 7756	AC004228	Genbank	4263838
Sequence 7757	AC004234	Genbank	2914672
Sequence 7758	AC004242	Genbank	3132845
Sequence 7759	AC004253	Genbank	3169206
Sequence 7760	AC004263	Genbank	2935616
Sequence 7761	AC004382	Genbank	3252819
Sequence 7762	AC004410	Genbank	2959558
Sequence 7763	AC004472	Genbank	2984582
Sequence 7764	AC004475	Genbank	2988396
Sequence 7765 Sequence 7766	AC004500	Genbank	2996639
Sequence 7767	AC004522 AC004526	Genbank Genbank	3006227
Sequence 7768	AC004526 AC004551	Genbank	3779035
Sequence 7769	AC004583	Genbank	3309005 3293210
Sequence 7770	AC004505	Genbank	3135391
Sequence 7771	AC004703	Genbank	3228509
Sequence 7772	AC004707	Genbank	3249127
Sequence 7773	AC004757	Genbank	3341707
Sequence 7774	AC004770	Genbank	3212836
Sequence 7775	AC004771	Genbank	3668108
Sequence 7776	AC004801	Genbank	4204244
Sequence 7777	AC004812	Genbank	3970970
Sequence 7778	AC004821	Genbank	4753291
Sequence 7779	AC004823	Genbank	4156207
Seguence 7780	AC004835	Genbank	4508154
Sequence 7781	AC004854	Genbank	4827328
Sequence 7782	AC004857	Genbank	3694665
Sequence 7783	AC004882	Genbank	5523837
Sequence 7784	AC004893	Genbank	3694662
Sequence 7785	AC004938	Genbank	3213059
Sequence 7786	AC004966	Genbank	5306307
Sequence 7787	AC004997	Genbank	5441941
Sequence 7788	AC005032	Genbank	5649384
Sequence 7789	AC005037	Genbank	4827310
Sequence 7790	AC005041	Genbank	4508118
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Sequence 7791         AC005058         Genbank         4156135           Sequence 7793         AC005071         Genbank         4508112           Sequence 7794         AC005081         Genbank         6560914           Sequence 7795         AC005088         Genbank         4753221           Sequence 7797         AC005192         Genbank         4218027           Sequence 7799         AC005214         Genbank         3282166           Sequence 7800         AC005225         Genbank         3282166           Sequence 7801         AC005318         Genbank         3282166           Sequence 7801         AC005318         Genbank         3283886           Sequence 7801         AC005318         Genbank         3365345           Sequence 7803         AC005324         Genbank         3365582           Sequence 7804         AC005339         Genbank         3367508           Sequence 7805         AC005366         Genbank         3367508           Sequence 7807         AC005392         Genbank         3337505           Sequence 7809         AC005480         Genbank         43367508           Sequence 7809         AC005480         Genbank         43367508           Seque				
Sequence 7793         AC005071         Genbank         6560914           Sequence 7794         AC005081         Genbank         6560914           Sequence 7795         AC005104         Genbank         4753221           Sequence 7797         AC005192         Genbank         3284765           Sequence 7799         AC005214         Genbank         3282166           Sequence 7800         AC005253         Genbank         328886           Sequence 7801         AC005318         Genbank         3385345           Sequence 7803         AC005324         Genbank         3385345           Sequence 7803         AC005325         Genbank         3365851           Sequence 7803         AC005325         Genbank         3365851           Sequence 7804         AC005339         Genbank         3367508           Sequence 7807         AC005369         Genbank         3367508           Sequence 7807         AC005392         Genbank         43367508           Sequence 7808         AC005480         Genbank         435818           Sequence 7809         AC005480         Genbank         4356194           Sequence 7809         AC005480         Genbank         4356194           Sequence	Sequence 7791	AC005042	Genbank	4156138
Sequence 7793         AC005071         Genbank         6560914           Sequence 7794         AC005081         Genbank         6560914           Sequence 7795         AC005104         Genbank         4753221           Sequence 7797         AC005192         Genbank         3284765           Sequence 7799         AC005214         Genbank         3282166           Sequence 7800         AC005253         Genbank         328886           Sequence 7801         AC005318         Genbank         3385345           Sequence 7803         AC005324         Genbank         3385345           Sequence 7803         AC005325         Genbank         3365851           Sequence 7803         AC005325         Genbank         3365851           Sequence 7804         AC005339         Genbank         3367508           Sequence 7807         AC005369         Genbank         3367508           Sequence 7807         AC005392         Genbank         43367508           Sequence 7808         AC005480         Genbank         435818           Sequence 7809         AC005480         Genbank         4356194           Sequence 7809         AC005480         Genbank         4356194           Sequence	Sequence 7792	AC005058	Genbank	4156135
Sequence 7794         AC005081         Genbank         6560914           Sequence 7795         AC005088         Genbank         4753221           Sequence 7798         AC005104         Genbank         3264575           Sequence 7799         AC005192         Genbank         3264575           Sequence 7780         AC005214         Genbank         3282166           Sequence 7800         AC005253         Genbank         3288886           Sequence 7801         AC005318         Genbank         3366582           Sequence 7802         AC005325         Genbank         3366582           Sequence 7804         AC005339         Genbank         3366581           Sequence 7805         AC005366         Genbank         3367505           Sequence 7806         AC005369         Genbank         3367505           Sequence 7809         AC005480         Genbank         3367505           Sequence 7810         AC005519         Genbank         5836194           Sequence 7811         AC005519         Genbank         5836194           Sequence 7812         AC005523         Genbank         580193           Sequence 7813         AC005536         Genbank         5901653           Sequence		AC005071	Genbank	4508112
Sequence 7795         AC005088         Genbank         4753221           Sequence 7796         AC005104         Genbank         4218027           Sequence 7797         AC005192         Genbank         3264575           Sequence 7799         AC005214         Genbank         3282166           Sequence 7800         AC005253         Genbank         3288866           Sequence 7801         AC005318         Genbank         3366582           Sequence 7803         AC005324         Genbank         3366581           Sequence 7804         AC005339         Genbank         3366581           Sequence 7806         AC005339         Genbank         3367508           Sequence 7806         AC005369         Genbank         3367508           Sequence 7807         AC005392         Genbank         3367508           Sequence 7808         AC005480         Genbank         4835818           Sequence 7810         AC005480         Genbank         5836194           Sequence 7811         AC005516         Genbank         5836194           Sequence 7811         AC005516         Genbank         5801651           Sequence 7813         AC005519         Genbank         4508129           Sequenc			Genbank	
Sequence 7796         AC005104         Genbank         3264575           Sequence 7797         AC005214         Genbank         3264575           Sequence 7798         AC005214         Genbank         3282166           Sequence 7799         AC005225         Genbank         6094661           Sequence 7801         AC005318         Genbank         3885345           Sequence 7802         AC005324         Genbank         3365581           Sequence 7803         AC005339         Genbank         3367508           Sequence 7805         AC005369         Genbank         3367508           Sequence 7806         AC005369         Genbank         3367508           Sequence 7807         AC005369         Genbank         3367508           Sequence 7808         AC005480         Genbank         3367508           Sequence 7810         AC005480         Genbank         4835818           Sequence 7811         AC005519         Genbank         5836194           Sequence 7811         AC005519         Genbank         3900835           Sequence 7811         AC005523         Genbank         3451032           Sequence 7814         AC005536         Genbank         3451032           Sequenc				
Sequence 7797         AC005192         Genbank         3264575           Sequence 7798         AC005214         Genbank         3282166           Sequence 7799         AC005225         Genbank         3094661           Sequence 7800         AC005253         Genbank         3288886           Sequence 7801         AC005318         Genbank         3366582           Sequence 7802         AC005325         Genbank         3366581           Sequence 7804         AC005339         Genbank         3367505           Sequence 7806         AC005366         Genbank         3367505           Sequence 7806         AC005369         Genbank         3367505           Sequence 7808         AC005480         Genbank         4835818           Sequence 7810         AC005480         Genbank         5836194           Sequence 7810         AC005516         Genbank         5836194           Sequence 7811         AC005519         Genbank         5836194           Sequence 7811         AC005516         Genbank         580198           Sequence 7813         AC005531         Genbank         4508129           Sequence 7814         AC005538         Genbank         4508129           Sequence				
Sequence 7798         AC005214         Genbank         3282166           Sequence 7799         AC005225         Genbank         6094661           Sequence 7800         AC005253         Genbank         328886           Sequence 7801         AC005318         Genbank         3366582           Sequence 7803         AC005325         Genbank         3366581           Sequence 7804         AC005339         Genbank         3367508           Sequence 7806         AC005369         Genbank         3367508           Sequence 7807         AC005392         Genbank         3367505           Sequence 7808         AC005480         Genbank         3396699           Sequence 7808         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         5836194           Sequence 7811         AC005516         Genbank         580163           Sequence 7811         AC005519         Genbank         580163           Sequence 7813         AC005523         Genbank         4508129           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005546         Genbank         3598724           Sequence 7			_ +	-
Sequence 7799         AC005225         Genbank         3288886           Sequence 7801         AC005318         Genbank         3288886           Sequence 7802         AC005318         Genbank         3366582           Sequence 7803         AC005325         Genbank         3366582           Sequence 7804         AC005339         Genbank         3367505           Sequence 7805         AC005369         Genbank         3367505           Sequence 7807         AC005392         Genbank         3397605           Sequence 7808         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         5836194           Sequence 7811         AC005516         Genbank         5836194           Sequence 7811         AC005519         Genbank         580183           Sequence 7811         AC005523         Genbank         4508129           Sequence 7813         AC005536         Genbank         4508129           Sequence 7814         AC005536         Genbank         4508129           Sequence 7815         AC005536         Genbank         4508129           Sequence 7816         AC005549         Genbank         3540153           Sequence				
Sequence 7800         AC005253         Genbank         3288886           Sequence 7801         AC005318         Genbank         3885345           Sequence 7802         AC005324         Genbank         3366582           Sequence 7803         AC005325         Genbank         3366581           Sequence 7804         AC005339         Genbank         3367508           Sequence 7806         AC005366         Genbank         3367505           Sequence 7807         AC005392         Genbank         3367505           Sequence 7808         AC005480         Genbank         3399669           Sequence 7809         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         5804985           Sequence 7811         AC005519         Genbank         5091653           Sequence 7812         AC005523         Genbank         4508129           Sequence 7814         AC005531         Genbank         4508129           Sequence 7814         AC005538         Genbank         450875           Sequence 7815         AC005549         Genbank         3478635           Sequence 7816         AC005612         Genbank         3540153           Sequence				
Sequence 7801         AC005318         Genbank         3885345           Sequence 7802         AC005324         Genbank         3366582           Sequence 7803         AC005325         Genbank         3365581           Sequence 7805         AC005366         Genbank         3367508           Sequence 7806         AC005369         Genbank         3367505           Sequence 7807         AC005392         Genbank         3395665           Sequence 7808         AC005480         Genbank         4835818           Sequence 7809         AC005480         Genbank         5836194           Sequence 7810         AC005516         Genbank         590835           Sequence 7811         AC005519         Genbank         5091653           Sequence 7812         AC005523         Genbank         4153875           Sequence 7814         AC005531         Genbank         4508129           Sequence 7815         AC005546         Genbank         3478635           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         35940153           Sequence 7818         AC005660         Genbank         3688099           Sequenc				
Sequence 7802         AC005324         Genbank         3366582           Sequence 7803         AC005325         Genbank         336581           Sequence 7804         AC005336         Genbank         3355454           Sequence 7806         AC005366         Genbank         3367508           Sequence 7807         AC005392         Genbank         3399669           Sequence 7808         AC005480         Genbank         4835818           Sequence 7809         AC005488         Genbank         5836194           Sequence 7810         AC005516         Genbank         590835           Sequence 7811         AC005519         Genbank         5091653           Sequence 7812         AC005523         Genbank         4508129           Sequence 7813         AC005531         Genbank         4508129           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005549         Genbank         3598724           Sequence 7816         AC005612         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7818         AC005612         Genbank         35980724           Sequence				
Sequence 7803         AC005325         Genbank         3366581           Sequence 7804         AC005339         Genbank         3355454           Sequence 7805         AC005369         Genbank         3367505           Sequence 7807         AC005392         Genbank         3399669           Sequence 7808         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         5836194           Sequence 7811         AC005519         Genbank         5901853           Sequence 7811         AC005519         Genbank         3900835           Sequence 7811         AC005519         Genbank         5091853           Sequence 7812         AC005523         Genbank         4508129           Sequence 7814         AC005531         Genbank         4508129           Sequence 7815         AC005546         Genbank         3598724           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7819         AC005629         Genbank         3598724           Sequence 7819         AC005670         Genbank         3608161           Sequenc	Seguence 7801	AC005318	Genbank	3885345
Sequence 7804         AC005339         Genbank         3355454           Sequence 7806         AC005369         Genbank         3367505           Sequence 7806         AC005369         Genbank         3367505           Sequence 7808         AC005392         Genbank         3399669           Sequence 7809         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         5900835           Sequence 7811         AC005516         Genbank         5091653           Sequence 7812         AC005523         Genbank         3451032           Sequence 7814         AC005531         Genbank         4508129           Sequence 7815         AC005538         Genbank         3478635           Sequence 7816         AC005546         Genbank         3478635           Sequence 7817         AC005612         Genbank         3540153           Sequence 7817         AC005612         Genbank         3540153           Sequence 7817         AC005629         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3688074           Sequenc	Sequence 7802	AC005324	Genbank	3366582
Sequence 7804         AC005339         Genbank         3355454           Sequence 7806         AC005369         Genbank         3367505           Sequence 7806         AC005369         Genbank         3367505           Sequence 7808         AC005392         Genbank         3399669           Sequence 7809         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         5900835           Sequence 7811         AC005516         Genbank         5091653           Sequence 7812         AC005523         Genbank         3451032           Sequence 7814         AC005531         Genbank         4508129           Sequence 7815         AC005538         Genbank         3478635           Sequence 7816         AC005546         Genbank         3478635           Sequence 7817         AC005612         Genbank         3540153           Sequence 7817         AC005612         Genbank         3540153           Sequence 7817         AC005629         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3688074           Sequenc	Sequence 7803	AC005325	Genbank	3366581
Sequence 7805         AC005366         Genbank         3367508           Sequence 7807         AC005369         Genbank         3367505           Sequence 7807         AC005392         Genbank         3399669           Sequence 7809         AC005480         Genbank         4835818           Sequence 7810         AC005516         Genbank         3900835           Sequence 7811         AC005519         Genbank         5091653           Sequence 7812         AC005523         Genbank         4153875           Sequence 7814         AC005531         Genbank         4508129           Sequence 7815         AC005534         Genbank         3478635           Sequence 7816         AC005549         Genbank         3540153           Sequence 7817         AC005549         Genbank         3540153           Sequence 7819         AC005629         Genbank         3540153           Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3698074           Sequence 7825         AC005789         Genbank         3789713           Sequenc		AC005339	Genbank	3355454
Sequence 7806         AC005369         Genbank         3367505           Sequence 7807         AC005392         Genbank         3399669           Sequence 7808         AC005480         Genbank         4835818           Sequence 7809         AC0055486         Genbank         5836194           Sequence 7811         AC005516         Genbank         5908351           Sequence 7812         AC005523         Genbank         3451032           Sequence 7814         AC005531         Genbank         4508129           Sequence 7814         AC005531         Genbank         4508129           Sequence 7815         AC005546         Genbank         3478635           Sequence 7815         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7818         AC005629         Genbank         3598724           Sequence 7821         AC0056670         Genbank         3688099           Sequence 7828         AC005686         Genbank         3608161           Sequence 7823         AC005768         Genbank         3688074           Sequence 7824         AC005789         Genbank         3698074           Seque		AC005366	Genbank	3367508
Sequence 7807         AC005392         Genbank         3399669           Sequence 7809         AC005480         Genbank         4835818           Sequence 7811         AC005516         Genbank         5836194           Sequence 7811         AC005516         Genbank         590835           Sequence 7812         AC005519         Genbank         5091653           Sequence 7813         AC005531         Genbank         4508129           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005538         Genbank         4508129           Sequence 7815         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7818         AC005629         Genbank         3598724           Sequence 7819         AC005629         Genbank         3588079           Sequence 7820         AC0056670         Genbank         3688099           Sequence 7821         AC005766         Genbank         3688074           Sequence 7823         AC005768         Genbank         3688074           Sequence 7824         AC005789         Genbank         3688074           Sequenc		•		
Sequence 7808         AC005480         Genbank         4835818           Sequence 7809         AC005516         Genbank         5836194           Sequence 7811         AC005516         Genbank         3900835           Sequence 7812         AC005523         Genbank         5091653           Sequence 7813         AC005531         Genbank         44508129           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005546         Genbank         3598724           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7818         AC005629         Genbank         3598724           Sequence 7819         AC005629         Genbank         3540153           Sequence 7819         AC005686         Genbank         3688099           Sequence 7821         AC005768         Genbank         3688074           Sequence 7821         AC005754         Genbank         3688074           Sequence 7823         AC0057589         Genbank         3702281           Sequence 7824         AC005899         Genbank         3789713           Seque				
Sequence 7809         AC005488         Genbank         5836194           Sequence 7810         AC005516         Genbank         3900835           Sequence 7811         AC005523         Genbank         5091653           Sequence 7812         AC005523         Genbank         3451032           Sequence 7813         AC005531         Genbank         4153875           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005549         Genbank         3478635           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7818         AC005629         Genbank         3540153           Sequence 7819         AC005629         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3688074           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         3688074           Sequence 7824         AC005789         Genbank         3702281           Sequenc				
Sequence 7810         AC005516         Genbank         390835           Sequence 7811         AC005519         Genbank         5091653           Sequence 7812         AC005523         Genbank         3451032           Sequence 7814         AC005531         Genbank         4153875           Sequence 7815         AC005546         Genbank         4508129           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3598724           Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC005629         Genbank         3540153           Sequence 7819         AC005629         Genbank         3688099           Sequence 7821         AC005686         Genbank         3688099           Sequence 7821         AC005726         Genbank         3688074           Sequence 7821         AC005768         Genbank         3688074           Sequence 7822         AC005768         Genbank         3688074           Sequence 7823         AC005789         Genbank         3789713           Sequence				
Sequence 7811         AC005519         Genbank         5091653           Sequence 7812         AC005523         Genbank         3451032           Sequence 7813         AC005531         Genbank         4153875           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005549         Genbank         3478635           Sequence 7816         AC005549         Genbank         3540153           Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC005629         Genbank         3540153           Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3688099           Sequence 7821         AC005726         Genbank         3688074           Sequence 7821         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         3598724           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7827         AC005841         Genbank         4731044           Sequenc				
Sequence 7812         AC005523         Genbank         3451032           Sequence 7813         AC005531         Genbank         4153875           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005546         Genbank         3478635           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC005629         Genbank         7243877           Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3688099           Sequence 7821         AC005726         Genbank         3688099           Sequence 7821         AC005726         Genbank         3688099           Sequence 7821         AC005768         Genbank         3688074           Sequence 7821         AC005768         Genbank         3688074           Sequence 7823         AC005789         Genbank         6598827           Sequence 7825         AC005828         Genbank         3702281           Sequence 7827         AC005829         Genbank         4731044           Sequenc				
Sequence 7813         AC005531         Genbank         4153875           Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005546         Genbank         3478635           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC0056829         Genbank         7243877           Sequence 7819         AC005686         Genbank         3688099           Sequence 7820         AC005686         Genbank         3688099           Sequence 7820         AC005686         Genbank         3688074           Sequence 7821         AC0057754         Genbank         3688074           Sequence 7823         AC005789         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3731044           Sequence 7827         AC005899         Genbank         4731044           Sequence 7828         AC005921         Genbank         4309927           Sequence 7830         AC005923         Genbank         4309927           Seque				
Sequence 7814         AC005538         Genbank         4508129           Sequence 7815         AC005548         Genbank         3478635           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC005629         Genbank         3688099           Sequence 7819         AC0056670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3688099           Sequence 7821         AC005726         Genbank         3688099           Sequence 7821         AC005754         Genbank         3688074           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005789         Genbank         6598827           Sequence 7824         AC005789         Genbank         3789713           Sequence 7826         AC005828         Genbank         4731044           Sequence 7827         AC005899         Genbank         4982556           Sequence 7828         AC005921         Genbank         4309922           Sequence 7830         AC005924         Genbank         4309922           Sequen				
Sequence 7815         AC005546         Genbank         3478635           Sequence 7816         AC005549         Genbank         3598724           Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC005629         Genbank         7243877           Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3688074           Sequence 7821         AC005726         Genbank         3688074           Sequence 7822         AC005754         Genbank         3688074           Sequence 7822         AC005768         Genbank         6598827           Sequence 7823         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005828         Genbank         4731044           Sequence 7827         AC005829         Genbank         4731044           Sequence 7828         AC005921         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309927           Sequence 7831         AC006026         Genbank         4156128           Sequenc				
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Sequence 7817         AC005612         Genbank         3540153           Sequence 7818         AC005629         Genbank         7243877           Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3810672           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         3688074           Sequence 7823         AC005768         Genbank         3688074           Sequence 7823         AC005789         Genbank         3688074           Sequence 7824         AC005789         Genbank         35827           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005828         Genbank         4731044           Sequence 7827         AC005899         Genbank         4982556           Sequence 7829         AC005921         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC006026         Genbank         4156128           Sequence				
Sequence 7818         AC005629         Genbank         7243877           Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3810672           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         4982556           Sequence 7828         AC005921         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309927           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         4156128           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4572650           Sequenc				
Sequence 7819         AC005670         Genbank         3688099           Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3810672           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005828         Genbank         4731044           Sequence 7827         AC005899         Genbank         4982556           Sequence 7828         AC005921         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         4314426           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4572650           Sequence 7835         AC006046         Genbank         4572650           Sequenc				
Sequence 7820         AC005686         Genbank         3608161           Sequence 7821         AC005726         Genbank         3810672           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         4982556           Sequence 7828         AC005921         Genbank         4309927           Sequence 7829         AC005923         Genbank         4309927           Sequence 7831         AC005984         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         4314426           Sequence 7833         AC006040         Genbank         4508120           Sequence 7835         AC006042         Genbank         4572650           Sequence 7836         AC006047         Genbank         4504701           Sequenc				
Sequence 7821         AC005726         Genbank         3810672           Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         4309926           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4572650           Sequence 7835         AC006042         Genbank         4572650           Sequence 7836         AC006046         Genbank         4204701           Sequenc				
Sequence 7822         AC005754         Genbank         3688074           Sequence 7823         AC005768         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         4982556           Sequence 7828         AC005921         Genbank         4309927           Sequence 7830         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006042         Genbank         4572650           Sequence 7836         AC006046         Genbank         4204701           Sequence 7838         AC006146         Genbank         4309313           Sequenc		AC005686	Genbank	
Sequence 7823         AC005768         Genbank         6598827           Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309927           Sequence 7831         AC005988         Genbank         4156128           Sequence 7831         AC005988         Genbank         6042101           Sequence 7833         AC006026         Genbank         4314426           Sequence 7834         AC006042         Genbank         4572650           Sequence 7835         AC006042         Genbank         4572650           Sequence 7836         AC006088         Genbank         4204701           Sequence 7839         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4309923           Sequenc		AC005726		
Sequence 7824         AC005789         Genbank         3702281           Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequenc	Sequence 7822	AC005754	Genbank	368 <b>807</b> 4
Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         4204701           Sequence 7837         AC006088         Genbank         4204701           Sequence 7839         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7842         AC006252         Genbank         4309923           Sequenc	Sequence 7823	AC005768	Genbank	6598827
Sequence 7825         AC005828         Genbank         3789713           Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4572650           Sequence 7835         AC006044         Genbank         4572650           Sequence 7836         AC006077         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7843         AC006255         Genbank         4309923           Sequenc	Sequence 7824	AC005789	Genbank	3702281
Sequence 7826         AC005841         Genbank         4731044           Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         4309923           Sequence 7843         AC006299         Genbank         4106997           Sequenc		AC005828	Genbank	3789 <b>713</b>
Sequence 7827         AC005899         Genbank         3935221           Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         4204701           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7844         AC006441         Genbank         4760435           Sequenc		AC005841	Genbank	4731044
Sequence 7828         AC005921         Genbank         4982556           Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         4204701           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7843         AC006299         Genbank         4106997           Sequence 7845         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666		AC005899	Genbank	3935221
Sequence 7829         AC005923         Genbank         4309927           Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7843         AC006299         Genbank         4106997           Sequence 7845         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666			Genbank	
Sequence 7830         AC005924         Genbank         4309926           Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7849         AC006165         Genbank         4160141           Sequence 7841         AC006241         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666			Genbank	
Sequence 7831         AC005988         Genbank         4156128           Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7843         AC006299         Genbank         4106997           Sequence 7845         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				4309926
Sequence 7832         AC006026         Genbank         6042101           Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7849         AC006165         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7833         AC006040         Genbank         4314426           Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7834         AC006042         Genbank         4508120           Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7835         AC006064         Genbank         4572650           Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7836         AC006077         Genbank         3935193           Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7837         AC006088         Genbank         4204701           Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7838         AC006146         Genbank         5836157           Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7839         AC006165         Genbank         3980464           Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7840         AC006241         Genbank         4160141           Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7841         AC006252         Genbank         4309923           Sequence 7842         AC006255         Genbank         5306223           Sequence 7843         AC006299         Genbank         4106997           Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
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Sequence 7844         AC006441         Genbank         4760435           Sequence 7845         AC006449         Genbank         6102666				
Sequence 7845 AC006449 Genbank 6102666				
Sequence 7846 AC006478 Genbank 4508145				
	Sequence 7846	AC006478	Genbank	4508145

Table 3-3

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Sequence 784		Genbank	6289252
Sequence 784		Genbank	4314418
Sequence 784		Genbank	4926863
Sequence 785		Genbank	5123989
Sequence 785	1 AC007022	Genbank	4753292
Sequence 785		Genbank	
Sequence 785:		Genbank	4371305
Sequence 7854		Genbank	4731066
Sequence 785			5903110
Sequence 7856		Genbank	6806840
		Genbank	6604532
Sequence 7857		Genbank	5306300
Sequence 7858		Genbank	6560920
Sequence 7859		Genbank	5230397
Sequence 7860		Genbank	5836156
Sequence 7861		Genbank	6466519
Sequence 7862		Genbank	5815499
Sequence 7863	AC007842	Genbank	5080755
Sequence 7864	AC007845	Genbank	7109495
Sequence 7865		Genbank	5649181
Sequence 7866		Genbank	
Sequence 7867	AC008134	Genbank	7025879
Sequence 7868	AC008990	_	5801655
Sequence 7869	AC009330	Genbank	7209394
Sequence 7870		Genbank	6553960
Sequence 7871	AC009514	Genbank	6016667
	AC010077	Genbank	5870275
Sequence 7872	AC011311	Genbank	6850247
Sequence 7873	AC012085	Genbank	6634755
Sequence 7874	AC016026	Genbank	7940357
Sequence 7875	AF000364	Genbank	2697102
Sequence 7876	AF001433	Genbank	2529704
Sequence 7877	AF001435	Genbank	2529708
Sequence 7878	AF001687	Genbank	2653735
Sequence 7879	AF001977	Genbank	2914738
Sequence 7880	AF002020	Genbank	2276462
Sequence 7881	AF002163	Genbank	2290769
Sequence 7882	AF002223	Genbank	4079670
Sequence 7883	AF002697	Genbank	2511528
Sequence 7884	AF002715	Genbank	2352276
Sequence 7885	AF002994	Genbank	2121297
Sequence 7886	AF002996	Genbank	2121299
Sequence 7887	AF004562	Genbank	3041872
Sequence 7888	AF005067	Genbank	6979018
Sequence 7889	AF005258	Genbank	2627428
Sequence 7890	AF005361	Genbank	2343115
Sequence 7891	AF006043	Genbank	2674061
Sequence 7892	AF006083	Genbank	2282031
Sequence 7893	AF006087	Genbank	
Sequence 7894	AF006515	Genbank	2282039
Sequence 7895	AF007170	Genbank	2645432
Sequence 7896	AF007544		2865251
Sequence 7897	AF007544 AF007791	Genbank	2970122
Sequence 7898	VEUU1121	Genbank	3779196
Sequence 7899	AF007871	Genbank	2358278
Sequence 7900	AF008303	Genbank	2580619
Sequence 7900	AF010187	Genbank	7025910
Sequence 7901	AF012872	Genbank	2326226
Sequence 7902	AF013970	Genbank	2801421

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Sequence 7903	AF014398	Genbank	8148061
Sequence 7904	AF015044	Genbank	4102712
Sequence 7905	AF015913	Genbank	2323409
Sequence 7906		Genbank	
	AF016050		2978559
Sequence 7907	AF016266	Genbank	2529 <b>5</b> 62
Sequence 7908	AF017790	Genbank	2501 <b>872</b>
Sequence 7909	AF020038	Genbank	3641397
Sequence 7910	AF020736	Genbank	3450954
	AF020833	Genbank	
Sequence 7911			2460199
Sequence 7912	AF021336	Genbank	2541972
Sequ <b>ence 7913</b>	AF021351	Genbank	2460207
Sequence 7914	AF022229	Genbank	2809382
Sequence 7915	AF023268	Genbank	2564910
Sequence 7916	AF023462	Genbank	2564670
Sequence 7917	AF023676	Genbank	3211721
Sequence 7918	AF025439	Genbank	2815605
Sequence 7919	AF025999	Genbank	2570853
Sequence 7920	AF026166	Genbank	4090928
Sequence 7921	AF026292	Genbank	25590 <b>0</b> 9
Sequence 7922	AF026548	Genbank	2583172
Sequence 7923	AF027299	Genbank	2739095
Sequence 7924	AF027302	Genbank	2522533
Sequence 7925	AF029308	Genbank	2564750
Sequence 7926	AF031416	Genbank	3213216
Sequence 7927	AF032456	Genbank	3004908
Sequence 7928	AF034373	Genbank	3820483
Sequence 7929	AF034756	Genbank	2654138
Sequence 7930	AF034759	Genbank	2653648
Sequence 7931	AF035191	Genbank	4104352
Sequence 7932	AF035286	Genbank	2661038
Sequence 7933	AF035295	Genbank	2661050
Sequence 7934	AF035296	Genbank	2661052
Sequence 7935	AF036536	Genbank	4104518
Sequence 7936	AF036682	Genbank	2707821
Sequence 7937	AF037448	Genbank	3037012
Sequence 7938	AF037643	Genbank	2746542
Sequence 7939	AF038196	Genbank	2795917
Sequence 7940	AF038318	Genbank	
			2708698
Sequence 7941	AF038391	Genbank	3123905
Sequence 7942	AF038392	Genbank	3123907
Sequence 7943	AF038554	Genbank	2895 <b>558</b>
Sequence 7944	AF039703	Genbank	2789655
Sequence 7945	AF040707	Genbank	3688794
Sequence 7946	AF040966	Genbank	2792367
Sequence 7947	AF041004	Genbank	4335800
•	AF041248	Genbank	3243117
Sequence 7948			
Sequence 7949	AF041259	Genbank	3335396
Sequence 7950	AF041381	Genbank	308 <b>0766</b>
Sequence 7951	AF041483	Genbank	3493528
Sequence 7952	AF042084	Genbank	2792517
Sequence 7953	AF042166	Genbank	3298596
Sequence 7954	AF042379	Genbank	2801700
Sequence 7955	AF043045	Genbank	3282770
	AF043045		
Sequence 7956		Genbank	3005062
Sequence 7957	AF044127	Genbank	4105189
Sequence 7958	AF044321	Genbank	3170263

Table 3-3

Sequence 7959	AF044588	Genbank	2865520
Sequence 7960	AF044773	Genbank	3002950
Sequence 7961	AF044954	Genbank	4164441
Sequence 7962	AF044956	Genbank	5326827
Sequence 7963	AF045606	Genbank	8277249
Sequence 7964	AF046025	Genbank	
	AF047181		4581063
Sequence 7965		Genbank	2909853
Sequence 7966	AF047184	Genbank	2909859
Sequence 7967	AF047431	Genbank	5514630
Sequence 7968	AF047432	Genbank	2911479
Sequence 7969	AF047437	Genbank	3335129
Sequence 7970	AF047442	Genbank	3335139
Sequence 7971	AF047472	Genbank	2921872
Sequence 7972	AF047599	Genbank	2906227
Sequence 7973	AF049910	Genbank	3435156
Sequence 7974	AF050078	Genbank	3818465
Sequence 7975	AF050641	Genbank	5326822
Sequence 7976	AF051782	Genbank	2947237
Sequence 7977	AF052059	Genbank	6851088
Sequence 7978	AF052106	Genbank	3360413
Sequence 7979	AF052130	Genbank	3360413
Sequence 7980	AF052155	Genbank	3360466
•	AF052178		
Sequence 7981 Sequence 7982		Genbank	3360489
	AF052179	Genbank	3360490
Sequence 7983	AF052182	Genbank	3360494
Sequence 7984	AF052186	Genbank	3360498
Sequence 7985	AF052504	Genbank	6650080
Sequence 7986	AF053069	Genbank	3136151
Sequence 7987	AF053074	Genbank	3283044
Sequence 7988	AF054187	Genbank	4092059
Sequence 7989	AF054284	Genbank	4033734
Sequence 7990	AF054990	Genbank	3005703
Sequence 7991	AF055014	Genbank	3005738
Sequence 7992	AF055017	Genbank	3005743
Sequence 7993	AF055022	Genbank	3005750
Sequence 7994	AF055029	Genbank	3005759
Sequence 7995	AF057160	Genbank	3694919
Sequence 7996	AF058392	Genbank	<b>30</b> 63679
Sequence 7997	AF058988	Genbank	3300089
Sequence 7998	AF059321	Genbank	3169563
Sequence 7999	AF061016	Genbank	3127126
Sequence 8000	AF061243	Genbank	3779243
Sequence 8001	AF061738	Genbank	4335940
Sequence 8002	AF061749	Genbank	3372676
Sequence 8003	AF062536	Genbank	3139076
Sequence 8004	AF063308	Genbank	4106355
Sequence 8005	AF064084	Genbank	3135668
Sequence 8006	AF064603	Genbank	3152659
Sequence 8007	AF065389	Genbank	3152702
Sequence 8008	AF067170	Genbank	4894373
Sequence 8009	AF067513	Genbank	3850305
Sequence 8010	AF067656	Genbank	3901271
Sequence 8011	AF067972	Genbank	4927369
Sequence 8012	AF068235	Genbank	4321975
Sequence 8013	AF068754	Genbank	
Sequence 8014	AF069301	Genbank	3283408
Ocquerice ou 14	VL009901	Genbank	3193335

Sequence 80	)15 A	F069984	Genbank	3242977
Sequence 80	016 A	F070550	Genbank .	3387913
Sequence 80		F070561	Genbank	3387928
Sequence 80		F070563	Genbank	3387931
Sequence 80		F070578	Genbank	3387950
Sequence 80		F070598	Genbank	3387976
Sequence 80		F070603	Genbank	3387983
Sequence 80		F070663	Genbank	4454701
Sequence 80		F071172	Genbank	5107833
Sequence 80		F071202	Genbank	
				3335172
Sequence 80		F071309	Genbank	3426319
Sequence 80		F071593	Genbank	3249712
Sequence 80		F071738	Genbank	4263875
Sequence 80		F071748	Genbank	3916211
Sequence 80		F072933	Genbank	5305428
Sequence 80		F073344	Genbank	5410229
Sequence 80		F074331	Genbank	5052074
Sequence 80		F075575	Genbank	3600027
Sequence 80		F076273	Genbank	4185718
Sequence 80		F076678	Genbank	4884560
Sequence 80	-	F076844	Genbank	4019216
Sequence 80		F077036	Genbank	4689119
Sequence 80	)37 A	F077 <b>042</b>	Genbank	4689131
Sequence 80	)38 A	F077188	Genbank	5565654
Sequence 80		F077203	Genbank	4679019
Sequence 80	40 A	F0773 <b>67</b>	Genbank	3820534
Sequence 80	141 A	F078103	Genbank	4218944
Sequence 80	42 A	F078107	Genbank	3342907
Sequence 80	43 A	F078671	Genbank	4877563
Sequence 80	144 A	F078776	Genbank	3511274
Sequence 80		F078845	Genbank	5531804
Sequence 80	46 A	F078847	Genbank	5531808
Sequence 80	47 A	F078854	Genbank	5531822
Sequence 80	48 A	F078863	Genbank	5531840
Sequence 80	49 A	F0 <b>78866</b>	Genbank	5531846
Sequence 80	50 A	F080561	Genbank	3746786
Sequence 80		F081484	Genbank	3420928
Sequence 80	52 A	F083068	Genbank	4808551
Sequence 80	53 A	F083106	Genbank	7555470
Sequence 80		F083190	Genbank	3599414
Sequence 80		F083246	Genbank	5106786
Sequence 80		F083255	Genbank	3435311
Sequence 80		F085355	Genbank	5114044
Sequence 80		F085357	Genbank	5114048
Sequence 80		F085359	Genbank	5114052
Sequence 80		F086234	Genbank	3483579
Sequence 80		F086280	Genbank	3483625
Sequence 80		F086406	Genbank	3483751
Sequence 80		F087438	Genbank	3820591
Sequence 80		F087986	Genbank	3523192
Sequence 80		F088060	Genbank	3523266
Sequence 80		F089841	Genbank	
Sequence 80		F090426	Genbank	4218954
Sequence 80		F090426 F090927		3982903
Sequence 80		F091035	Genbank	6690220
Sequence 80		_	Genbank	6002584
ocdacure 90	O A	F0 <b>9107</b> 3	Genbank	3859983

Table 3-3

Sequence 8071	AF092136	Genbank	5138921
Sequence 8072	AF092565	Genbank	3661609
Sequence 8073	AF093239	Genbank	3661615
Sequence 8074	AF094516	Genbank	3820613
Sequence 8075	AF095192	Genbank	4322819
Sequence 8076	AF095448	Genbank	4063889
Sequence 8077	AF095890	Genbank	4185790
Sequence 8078	AF096160	Genbank	4454849
Sequence 8079	AF097514	Genbank	4808600
Sequence 8080	AF098865	Genbank	4204674
Sequence 8081	AF098948	Genbank	5456746
Sequence 8082	AF100615	Genbank	4808624
Sequence 8083	AF100755	Genbank	5410295
Sequence 8084	AF100756	Genbank	
Sequence 8085	AF103800	Genbank	5410297
	AF103600 AF104913	Genbank	6048967
Sequence 8086	AF104913 AF105366		3941723
Sequence 8087		Genbank	5106522
Sequence 8088	AF106798	Genbank	5805207
Sequence 8089	AF107405	Genbank	5531903
Sequence 8090	AF107406	Genbank	5531905
Sequence 8091	AF110184	Genbank	5565858
Sequence 8092	AF111168	Genbank	4186181
Sequence 8093	AF111853	Genbank	6560625
Sequence 8094	AF112227	Genbank	4545218
Sequence 8095	AF113019	Genbank	6642761
Sequence 8096	AF113129	Genbank	6523820
Sequence 8097	AF113132	Genbank	5326801
Sequence 8098	AF113249	Genbank	6650673
Sequence 8099	AF113251	Genbank	5852417
Sequence 8100	AF113702	Genbank	6855636
Sequence 8101	AF116241	Genbank	5006438
Sequence 8102	AF117888	Genbank	5732617
Sequence 8103	AF118124	Genbank	4235636
Sequence 8104	AF124145	Genbank	5931954
Sequence 8105	AF125102	Genbank	5106999
Sequence 8106	AF125525	Genbank	4689281
Sequence 8107	AF126021	Genbank	6563273
Sequence 8108	AF126246	Genbank	4406523
Sequence 8109	AF126782	Genbank	6318547
Sequence 8110	AF126799	Genbank	4406527
Sequence 8111	AF127138	Genbank	5922724
Sequence 8112	AF127563	Genbank	4455124
Sequence 8113	AF127761	Genbank	4455128
Sequence 8114	AF127966	Genbank	5532271
Sequence 8115	AF128458	Genbank	7140573
Sequence 8116	AF129930	Genbank	4558509
Sequence 8117	AF131742	Genbank	4406554
Sequence 8118	AF131797	Genbank	4406625
Sequence 8119	AF131802	Genbank	4406633
Sequence 8120	AF131820	Genbank	4406655
Sequence 8121	AF131826	Genbank	4406662
Sequence 8122	AF132033	Genbank	5524202
Sequence 8123	AF132856	Genbank	4809025
Sequence 8124	AF132940	Genbank	4680650
Sequence 8125	AF132946	Genbank	4680 <b>662</b>
Sequence 8126	AF132965	Genbank	4680700

Table 3-3

Sequence 8127	AF134726	Genbank	4529886
Sequence 8128	AF134803	Genbank	4868362
Sequence 8129	AF136271	Genbank	4732145
Sequence 8130	AF136715	Genbank	5019773
Sequence 8131	AF140710	Genbank	6073848
Sequence 8132	AF141347	Genbank	4929133
Sequence 8133	AF141870	Genbank	5006601
Sequence 8134	AF143536	Genbank	4929219
Sequence 8135	AF145732	Genbank	5579330
Sequence 8136	AF147271	Genbank	5823152
Sequence 8137	AF147331	Genbank	4761682
Sequence 8138	AF147410	Genbank	4761761
Sequence 8139	AF147787	Genbank	5231122
Sequence 8140	AF148513	Genbank	4809268
Sequence 8141	AF149045	Genbank	6010479
Sequence 8142	AF150087	Genbank	5107162
Sequence 8143	AF151018	Genbank	7106757
Sequence 8144	AF151020	Genbank	7106761
Sequence 8145	AF151042	Genbank	7106805
Sequence 8146	AF151051	Genbank	7106823
Sequence 8147	AF151109	Genbank	6166337
Sequence 8148	AF151799	Genbank	4929550
Sequence 8149	AF151799	Genbank	4929562
•	AF151808	Genbank	4929568
Sequence 8150			
Sequence 8151	AF151811	Genbank	4929574
Sequence 8152	AF151826	Genbank	4929604
Sequence 8153	AF151841	Genbank	4929634
Sequence 8154	AF151846	Genbank	4929644
Sequence 8155	AF151861	Genbank	4929674
Sequence 8156	AF151869	Genbank	4929690
Sequence 8157	AF151886	Genbank	4929724
Sequence 8158	AF151902	Genbank	4929756
Sequence 8159	AF151904	Genbank	4929760
Sequence 8160	AF152961	Genbank	5499740
Sequence 8161	AF155095	Genbank	5360084
Sequence 8162	AF155110	Genbank	5360114
Sequence 8163	AF155238	Genbank	9186822
Sequence 8164	AF156165	Genbank	7145093
Sequence 8165	AF157378	Genbank	5714631
Sequence 8166	AF159295	Genbank	5714635
Sequence 8167	AF161371	Genbank	6841155
Sequence 8168	AF161417	Genbank	6841247
Sequence 8169	AF161448	Genbank	6841309
Sequence 8170	AF161455	Genbank	6841323
Sequence 8171	AF161458	Genbank	6841439
Sequence 8172	AF161485	Genbank	6841493
Sequence 8173	AF161491	Genbank	6841505
Sequence 8174	AF161493	Genbank	6841509
Sequence 8175	AF161511	Genbank	6841545
Sequence 8176		Genbank	
Sequence 8177	AF161530	Genbank	6841583
	AF163441		6969979
Sequence 8178	AF165967	Genbank	5802826
Sequence 8179	AF167572	Genbank	6164703
Sequence 8180	AF167706	Genbank	6979310
Sequence 8181	AF168787	Genbank	7239175
Sequence 8182	AF168956	Genbank	5702387

Table 3-3

Sequence 8183	AF180682	Genbank	6684524
Sequence 8184	AF181265	Genbank	7212810
Sequence 8185	AF182316	Genbank	6731234
Sequence 8186	AF187320	Genbank	6164847
Sequence 8187	AF187554	Genbank	6653225
Sequence 8188	AF190167	Genbank	
Sequence 8189	AF190725		6456117
Sequence 8190		Genbank	6118538
	AF191339	Genbank	6180012
Sequence 8191	AF191340	Genbank	6180014
Sequence 8192	AF192968	Genbank	6449082
Sequence 8193	AF196779	Genbank	61 <b>801</b> 70
Sequence 8194	AF196969	Genbank	6 <b>2890</b> 73
Sequence 8195	AF198444	Genbank	7025338
Sequence 8196	AF198487	Genbank	6941955
Sequence 8197	AF199596	Genbank	<b>684204</b> 9
Sequence 8198	AF203815	Genbank	6979641
Sequence 8199	AF203978	Genbank	6503203
Sequence 8200	AF207907	Genbank	665 <b>36</b> 48
Sequence 8201	AF212162	Genbank	6665785
Sequence 8202	AF220417	Genbank	7678700
Sequence 8203	AF223408	Genbank	6911266
Sequence 8204	AF224669	Genbank	7012905
Sequence 8205	AF227218	Genbank	7159279
Sequence 8206	AJ000186	Genbank	2463195
Sequence 8207	AJ000332	Genbank	2274967
Sequence 8208	AJ000644	Genbank	2695707
Sequence 8209	AJ001050	Genbank	2832345
Sequence 8210	AJ001443	Genbank	6006514
Sequence 8211	AJ002955	Genbank	2632122
Sequence 8212	AJ004832	Genbank	
Sequence 8213	AJ006291	Genbank	2982500 3805937
Sequence 8214	AJ007398	Genbank	
Sequence 8215	AJ007509	Genbank	3668140
Sequence 8216	AJ007670		3319955
Sequence 8217	AJ010597	Genbank	3560123
Sequence 8218		Genbank	3559873
Sequence 8219	AJ011930	Genbank	3859769
	AJ012463	Genbank	3850049
Sequence 8220	AJ012499	Genbank	5441359
Sequence 8221	AJ131186	Genbank	5689737
Sequence 8222	AJ223366	Genbank	6691913
Sequence 8223	AJ224442	Genbank	2911586
Sequence 8224	AJ245416	Genbank	5701851
Sequence 8225	AJ250475	Genbank	8919097
Sequence 8226	AJ275986	Genbank	7228110
Sequence 8227	AK000056	Genbank	7019892
Sequence 8228	AK000161	Genbank	7020067
Sequence 8229	AK000181	Genbank	7020098
Sequence 8230	AK000221	Genbank	7020163
Sequence 8231	AK000228	Genbank	7020174
Sequence 8232	AK000260	Genbank	7020221
Sequence 8233	AK000284	Genbank	7020260
Sequence 8234	AK000286	Genbank	7020264
Sequence 8235	AK000298	Genbank	7020286
Sequence 8236	AK000301	Genbank	7020291
Sequence 8237	AK000357	Genbank	7020388
Sequence 8238	AK000361	Genbank	7020395
			, 520030

Sequence 8239	AK000371	Genbank	7020413
Sequence 8240	AK000373	Genbank	7020417
Sequence 8241	AK000398	Genbank	7020459
Sequence 8242	AK000407	Genbank	7020474
Sequence 8243	AK000436	Genbank	7020522
Sequence 8244	AK000474	Genbank	7020586
Sequence 8245	AK000498	Genbank	7020628
Sequence 8246	AK000505	Genbank	7020640
Sequence 8247	AK000543	Genbank	7020709
Sequence 8248	AK000544	Genbank	7020710
Sequence 8249	AK000558	Genbank	7020734
Sequence 8250	AK000560	Genbank	7020738
Sequence 8251	AK000595	Genbank	7020796
Sequence 8252	AK000595	Genbank	7020730
	AK000651	Genbank	7020882
Sequence 8253	AK000664	Genbank	7020002
Sequence 8254		Genbank	7020901
Sequence 8255	AK000669		
Sequence 8256	AK000714	Genbank	7020975
Sequence 8257	AK000739	Genbank	7021014
Sequence 8258	AK000740	Genbank	7021016
Sequence 8259	AK000745	Genbank	7021025
Sequence 8260	AK000757	Genbank	7021043
Sequence 8261	AK000759	Genbank	7021046
Sequence 8262	AK000799	Genbank	7021103
Sequence 8263	AK000859	Genbank	7021183
Sequence 8264	AK000946	Genbank	7021928
Sequence 8265	AK000982	Genbank	7021980
Sequence 8266	AK000990	Genbank	7021992
Sequence 8267	AK001090	Genbank	7022139
Sequence 8268	AK001112	Genbank	7022170
Sequence 8269	AK001123	Genbank	7022186
Sequence 8270	AK001184	Genbank	7022279
Sequence 8271	AK001188	Genbank	7022286
Sequence 8272	AK001262	Genbank	7022406
Sequence 8273	AK001264	Genbank	7022409
Sequence 8274	AK001269	Genbank	7022419
Sequence 8275	AK001277	Genbank	7022431
Sequence 8276	AK001281	Genbank	7022437
Sequence 8277	AK001288	Genbank	7022448
Sequence 8278	AK001292	Genbank	7022455
Sequence 8279	AK001313	Genbank	7022490
Sequence 8280	AK001330	Genbank	7022520
Sequence 8281	AK001344	Genbank	7022543
Sequence 8282	AK001363	Genbank	7022575
Sequence 8283	AK001364	Genbank	7022577
Seguence 8284	AK001366	Genbank	7022579
Sequence 8285	AK001387	Genbank	7022615
Sequence 8286	AK001392	Genbank	7022621
Sequence 8287	AK001395	Genbank	7022625
Sequence 8288	AK001333	Genbank	7022649
Sequence 8289	AK001410 AK001436		7022692
	AK001436 AK001478	Genbank	
Sequence 8290	AK001478 AK001492	Genbank	7022760 7022781
Sequence 8291		Genbank	
Sequence 8292	AK001508	Genbank	7022805
Sequence 8293	AK001521	Genbank	7022828
Sequence 8294	AK001539	Genbank	7022856

Table 3-3

Sequence 8295	AK001565	Genbank	7022895
Sequence 8296	AK001601	Genbank	7022955
Sequence 8297	AK001613	Genbank	7022975
Sequence 8298	AK001627	Genbank	7022996
Sequence 8299	AK001636	Genbank	7023010
Sequence 8300	AK001675	Genbank	7023079
Sequence 8301	AK001690	Genbank	7023105
Sequence 8302	AK001691	Genbank	7023106
Sequence 8303	AK001700	Genbank	7023122
Sequence 8304	AK001714	Genbank	7023146
Sequence 8305	AK001716	Genbank	7023150
Sequence 8306	AK001725	Genbank	7023165
Sequence 8307	AK001741	Genbank	7023192
Sequence 8308	AK001775	Genbank	7023255
Sequence 8309	AK001829	Genbank	7023339
Sequence 8310	AK001852	Genbank	7023376
Sequence 8311	AK001939	Genbank	7023513
Sequence 8312	AK001962	Genbank	7023551
Sequence 8313	AK002019	Genbank	7023647
Sequence 8314	AK002051	Genbank	7023698
Sequence 8315	AK002060	Genbank	7023713
Sequence 8316	AK002094	Genbank	7023767
Sequence 8317	AK002136	Genbank	7023830
Sequence 8318	AK002165	Genbank	7023876
Sequence 8319	AK002212	Genbank	7023953
Sequence 8320	AL008639	Genbank	6599068
Sequence 8321	AL008721	Genbank	3171883
Sequence 8322	AL020995	Genbank	5881341
Sequence 8323	AL021453	Genbank	3413288
Sequence 8324	AL021546	Genbank	2826890
Sequence 8325	AL021707	Genbank	4582132
Sequence 8326	AL021807	Genbank	9367202
Sequence 8327	AL021917	Genbank	9367203
Sequence 8328	AL021978	Genbank	3135963
Sequence 8329	AL022157	Genbank	3319676
Sequence 8330	AL022313	Genbank	4200326
Sequence 8331	AL022318	Genbank	4826439
Sequence 8332	AL022339	Genbank	3355600
Sequence 8333 Sequence 8334	AL022394	Genbank	9581758
Sequence 8335	AL023803 AL024498	Genbank	9408726
Sequence 8336	AL024498 AL031387	Genbank	5924006
Sequence 8337	AL031427	Genbank Genbank	4493578
Sequence 8338	AL031427 AL031680	Genbank Genbank	4835258
Sequence 8339	AL031685	Genbank Genbank	9368417
Sequence 8340	AL031733	Genbank	9368423
Sequence 8341	AL031733 AL031778	_	4826495
Sequence 8342	AL034743	Genbank Genbank	4153958
Sequence 8343	AL034379	Genbank	5931807
Sequence 8344	AL034379 AL034380	Genbank	5918013 7981257
Sequence 8345	AL035420	Genbank	5918161
Sequence 8346	AL035551	Genbank	4826429
Sequence 8347	AL035587	Genbank	6002306
Sequence 8348	AL035659	Genbank	6562059
Sequence 8349	AL035683	Genbank	7288039
Sequence 8350	AL049219	Genbank	4499949
		-0.104111	7700070

Sequence 8351	AL049367	Genbank	4500158
Sequence 8352	AL049539	Genbank	6456855
Sequence 8353	AL049546	Genbank	4678775
Sequence 8354	AL049589	Genbank	5679567
Sequence 8355	AL049611	Genbank	5881343
Sequence 8358	AL049646	Genbank	9368462
Sequence 8357	AL049712	Genbank	5629919
Sequence 8358	AL049713	Genbank	6006528
Sequence 8359	AL049742	Genbank	5419783
Sequence 8360	AL049795	Genbank	6010175
Sequence 8361	AL049801	Genbank	4902677
Sequence 8362	AL049821	Genbank	5791513
Sequence 8363	AL049943	Genbank	4884187
Sequence 8364	AL050011	Genbank	4884080
Sequence 8365	AL050028	Genbank	4884267
Sequence 8366	AL050107	Genbank	4884135
Sequence 8367	AL050118	Genbank	4884143
Sequence 8368	AL050135	Genbank	4884344
Sequence 8369	AL050187	Genbank	4884402
Sequence 8370	AL050208	Genbank	4884447
Sequence 8371	AL050274	Genbank	4886452
Sequence 8372	AL050332	Genbank	6010176
Sequence 8373	AL050367	Genbank	4914600
Sequence 8374	AL050396	Genbank	4914618
Sequence 8375	AL078581	Genbank	6967288
Sequence 8376	AL078594	Genbank	7161750
Sequence 8377	AL078611	Genbank	5019312
Sequence 8378	AL078621	Genbank	6013067
Sequence 8379	AL079298	Genbank	5102736
Sequence 8380	AL079352	Genbank	7799 <b>779</b>
Sequence 8381	AL080080	Genbank	5262491
Sequence 8382	AL080155	Genbank	5262612
Sequence 8383	AL080157	Genbank	5262616
Sequence 8384	AL080172	Genbank	5262642
Sequence 8385	AL096738	Genbank	5419870
Sequence 8386	AL096740	Genbank	5419896
Sequence 8387	AL096752	Genbank	5419888
Sequence 8388	AL096791	Genbank	6066332
Sequence 8389	AL109788	Genbank	5725476
Sequence 8390	AL109794	Genbank	5725481
Sequence 8391	AL109925	Genbank	7105935
Sequence 8392	AL110244	Genbank	5817191
Sequence 8393	AL110268	Genbank	5817041
Sequence 8394	AL110273	Genbank	5817091
Sequence 8395	AL117398	Genbank	5911918
Sequence 8396	AL117409	Gen <b>bank</b>	5912046
Sequence 8397	AL117429	Gen <b>ban</b> k	5911863
Sequence 8398	AL117482	Genbank	5911960
Sequence 8399	AL117500	Genbank	5912005
Sequence 8400	AL117572	Genbank	5912124
Sequence 8401	AL117573	Genbank	5912125
Sequence 8402	AL117579	Genbank	5912135
Sequence 8403	AL117594	Genbank	5912158
Sequence 8404	AL117607	Genbank	5912177
Sequence 8405	AL117694	Genbank	8248721
Sequence 8406	AL121575	Genbank	7264053

Sequence 8407	AL121790	Genbank	8919824
Sequence 8408	AL122020	Genbank	8176920
Sequence 8409	AL133036	Genbank	6453441
Sequence 8410	AL133064	Genbank	6453490
Sequence 8411	AL133078	Genbank	6453522
Sequence 8412	AL133215	Genbank	7228177
Sequence 8413	AL133246	Genbank	7594587
Sequence 8414	AL133353	Genbank	6706037
Sequence 8415	AL133455	Genbank	7019734
Sequence 8416	AL133555	Genbank	6599122
Sequence 8417	AL133583	Genbank	6599167
Sequence 8418	AL133589	Genbank	6599176
Sequence 8419	AL133609	Genbank	6599219
Sequence 8420	AL135744	Genbank	6682291
Sequence 8421	AL135858	Genbank	7159622
Sequence 8422	AL136295	Genbank	6850939
Sequence 8423	AL136543	Genbank	6807646
Sequence 8424	AL137263	Genbank	6807692
Sequence 8425	AL137304	Genbank	6807768
Sequence 8426	AL137327	Genbank	6807818
Sequence 8427	AL137406	Genbank	6807955
Sequence 8428	AL137471	Genbank	6808079
Sequence 8429	AL137540	Genbank	6808218
Sequence 8430	AL137548	Genbank	6808231
Sequence 8431	AL137585	Genbank	6808304
Sequence 8432	AL137641	Genbank	6808443
Sequence 8433	AL137681	Genbank	6807931
Sequence 8434	AL137683	Genbank	6807935
Sequence 8435	AL137727	Genbank	6808256
Sequence 8436	AL137743	Genbank	6808323
Sequence 8437	AL139165	Genbank	6969146
Sequence 8438	AP000010	Genbank	4666256
Sequence 8439	AP000d46	Genbank	3132356
Sequence 8440	AP000066	Genbank	4579987
Sequence 8441	AP000493	Genbank	5926660
Sequence 8442	AP000501	Genbank	5926688
Sequence 8443	AP000529	Genbank	5931507
Sequence 8444	AP000694	Genbank	6693626
Sequence 8445	AP001115	Genbank	6899854
Sequence 8446	AP001137	Genbank	6970361
Sequence 8447	D00591	Genbank	220051
Sequence 8448	D01038	Genbank	220140
Sequence 8449	D10040	Genbank	219899
Sequence 8450	D10495	Genbank	520586
Sequence 8451	D12981	Genbank	219979
Sequence 8452	D13119	Genbank	285909
Sequence 8453	D13286	Genbank	496368
Sequence 8454	D13540	Genbank	220071
Sequence 8455	D13627	Genbank	286010
Sequence 8456	D13635	Genbank	285982
Sequence 8457	D13748	Genbank	219402
Sequence 8458	D14662	Genbank	285948
Sequence 8459	D14689	Genbank	285956
Sequence 8460	D14694	Genbank	603801
Sequence 8461	D14811	Genbank	285966
Sequence 8462	D16307	Genbank	303594

Table 3-3

Sequence 8463	D16562	Genbank	506336
Sequence 8464	D21063	Genbank	434752
Sequence 8465	D21163	Genbank	434758
Sequence 8466	D21235	Genbank	498145
Sequence 8467	D21254	Genbank	575577
Sequence 8468	D21261	Genbank	434762
Sequence 8469	D21262	Genbank	434764
Sequence 8470	D21852	Genbank	434768
Sequence 8471	D25215	Genbank	517114
Sequence 8472	D26156	Genbank	505087
Sequence 8473	D26181	Genbank	473712
Sequence 8474	D26351	Genbank	450470
Sequence 8475	D26485	Genbank	468934
Sequence 8476	D28473	Genbank	551621
Sequence 8477	D28476	Genbank	460710
Sequence 8478	D28589	Genbank	460714
		Genbank	473940
Sequence 8479	D29954	Genbank	473948
Sequence 8480	D29958		
Sequence 8481	D30756	Genbank	488500
Sequence 8482	D31762	Genbank	498149
Sequence 8483	D31764	Genbank	498153
Sequence 8484	D31767	Genbank	505091
Sequence 8485	D31885	Genbank	505097
Sequence 8486	D31888	Genbank	506340
Sequence 8487	D37766	Genbank	2429078
Sequence 8488	D38047	Genbank	1037163
Sequence 8489	D38048	Genbank	1531532
Sequence 8490	D38 <b>073</b>	Genbank	862331
Sequence 8491	D38522	Genbank	559331
Sequence 8492	D38 <b>524</b>	Genbank	633070
Sequence 8493	D38549	Genbank	559702
Sequence 8494	D385 <b>53</b>	Genbank	559714
Sequence 8495	D38555	Genbank	559716
Sequence 8496	D38616	Genbank	1304117
Sequence 8497	D42047	Genbank	577306
Sequence 8498	D42055	Genbank	5 <b>773</b> 12
Sequence 8499	D42085	Genbank	577316
Sequence 8500	D43682	Genbank	1060913
Sequence 8501	D43949	Genbank	603952
Sequence 8502	D43950	Genbank	603954
Sequence 8503	D43951	Genbank	603956
Sequence 8504	D45021	Genbank	624872
Sequence 8505	D45131	Genbank	1304103
Sequence 8506	D49489	Genbank	1136742
Sequence 8507	D49490	Genbank	1072306
Sequence 8508	D50369	Genbank	2605589
Sequence 8509	D50372	Genbank	2605593
Sequence 8510	D50923	Genbank	1469188
Sequence 8511	D50929	Genbank	1469200
Sequence 8512	D55716	Genbank	1255616
Sequence 8513	D63475	Genbank	1665724
Sequence 8514	D63478	Genbank	1469869
Sequence 8515	D63479	Genbank	6633998
Sequence 8516	D63486	Genbank	1469885
Sequence 8517	D63780	Genbank	2196444
Sequence 8518	D63861	Genbank	1769811
	20001	Combanic	.,05511

Table 3-3

Sequence 8519	D63875	Genbank	961441
Sequence 8520	D78014	Genbank	1330241
Sequence 8521	D79205	Genbank	1754620
Sequence 8522	D79984	Genbank	1136385
Sequence 8523	D79991	Genbank	1136397
Sequence 8524	D80000	Genbank	1136415
Sequence 8525	D80009	Genbank	1136433
Sequence 8526	D80010	Genbank	1136435
Sequence 8527	D82348	Genbank	1311461
Sequence 8528	D83077	Genbank	1304131
Sequence 8529	D83778	Genbank	1228038
Sequence 8530	D83781	Genbank	1228044
Sequence 8531	D83782	Genbank	1228046
Sequence 8532	D84064	Genbank	2618587
Sequence 8533	D86198	Genbank	3062805
Sequence 8534	D86962	Genbank	1503997
Sequence 8535	D86965	Genbank	
Sequence 8536	D86977	Genbank	1504003
Sequence 8537	D86978	Genbank	1504027
Sequence 8538	D86985	Genbank	1504029
Sequence 8539	D86987	Genbank	6634002
Sequence 8540	D80987	Genbank	3869186
Sequence 8541	D87436		1817551
Sequence 8542	D87447	Genbank	1665766
Sequence 8543	D87452	Genbank Genbank	1665782
Sequence 8544	D87464	Genbank	1665792
Sequence 8545	D87470	Genbank	1665812
Sequence 8546		Genbank	1665822
Sequence 8547	D87717 D87735	Genbank Genbank	1663709
Sequence 8548	D88674	Genbank Genbank	1620021
Sequence 8549	D88687	Genbank	2641951
Sequence 8550	D89980	Genbank	1843433
Sequence 8551	D90391	Genbank	2804272
Sequence 8552	E01094	Genbank	219491
Sequence 8553	E01787	Genbank	2169353
Sequence 8554	E01816	Genbank	2170040
Sequence 8555	E01956	Genbank	2170068
Sequence 8556	E02164	Genbank	2170204 2170402
Sequence 8557	E02628	Genbank	2170856
Sequence 8558	E06721	Genbank	2174903
Sequence 8559	E12458	Genbank	3251291
Sequence 8560	E13330	Genbank	3252135
Sequence 8561	E14580	Genbank	
Sequence 8562	E14872	Genbank	5709263
Sequence 8563	E15239	Genbank	5709555
Sequence 8564	E15695		5709922 5710378
Sequence 8565	E17361	Genbank Genbank	5710378 5712044
Sequence 8566	J02642	Genbank	5712044
Sequence 8567	J02042 J02923	Genbank	182862
Sequence 8568	J03007		189501
Sequence 8569	J03007	Genbank	179211
Sequence 8570	J03202	Genbank Genbank	337755
Sequence 8571	J03202 J03473	Genbank Genbank	186916
Sequence 8572	J03473 J03518		337423
Sequence 8573	J03518 J03528	Genbank	340389
Sequence 8574	J03526 J03575	Genbank	188671
	303375	Genbank	189737

Table 3-3

Sequence 8575	J03576	Genbank	189753
Sequence 8576	J03592	Genbank	339722
Sequence 8577	J03799	Genbank	186840
Sequence 8578	J03827	Genbank	340418
Sequence 8579	J04031	Genbank	187464
Sequence 8580	J04088	Genbank	292829
Sequence 8581	J04164	Genbank	177801
Sequence 8582	J04173	Genbank	551173
Sequence 8583	J04444	Genbank	181239
Sequence 8584	J04543	Genbank	338243
Sequence 8585	J04611	Genbank	178649
Sequence 8586	J04615	Genbank	338246
Sequence 8587	J04988	Genbank	184422
Sequence 8588	J05594	Genbank	1203981
Sequence 8589	J05633	Genbank	186504
Sequence 8590	K00558	Genbank	340020
Sequence 8591	L00160	Genbank	189904
Sequence 8592	L00635	Genbank	292032
Sequence 8593	L01457	Genbank	179282
Sequence 8594	L04733	Genbank	307084
Sequence 8595	L05093	Genbank	401844
Sequence 8596	L05425	Genbank	179284
Sequence 8597	L06237	Genbank	437000
Sequence 8598	L06328	Genbank	340200
Sequence 8599	L06419	Genbank	190073
Sequence 8600	L07044	Genbank	291894
Sequence 8601	L08238	Genbank	307198
Sequence 8602	L09159	Genbank	307374
Sequence 8603	L09561	Genbank	3192937
Sequence 8604	L11931	Genbank	307421
Sequence 8605	L11932	Genbank	307423
Sequence 8606	L13850	Genbank	292165
Sequence 8607	L14076	Genbank	307437
Sequence 8608	L14283	Genbank	307355
Sequence 8609	L14599	Genbank	348238
Sequence 8610	L14837	Genbank	292937
Sequence 8611	L16783	Genbank	452438
Sequence 8612	L19597	Genbank	306467
Sequence 8613	L19956	Genbank	306456
Sequence 8614	L20941	Genbank	507251
Sequence 8615	1.22005	Genbank	388308
Sequence 8616	L22009	Genbank	347313
Sequence 8617	L22740	Genbank	409222
Sequence 8618	L23959	Genbank	414316
Sequence 8619	L25270	Genbank	457136
Sequence 8620	L25876	Genbank	414584
Sequence 8621	L25880	Genbank	450272
Sequence 8622	L26524	Genbank	457232
Sequence 8623	L27211	Genbank	558656
Sequence 8624	L27711	Genbank	808006
Sequence 8625	L29216	Genbank	632965
Sequence 8626	L29296	Genbank	559042
Sequence 8627	L29555	Genbank	522196
Sequence 8628	L34075	Genbank	508481
Sequence 8629	L36056	Genbank	561631
Sequence 8630	L36818	Genbank	556191

Table 3-3

Sequence 8631	L38961	Genbank	624703
Sequence 8632	L39068	Genbank	994714
Sequence 8633	L39210	Genbank	1702963
Sequence 8634	L39211	Genbank	755645
Sequence 8635	L40326	Genbank	695361
Sequence 8636	L40586	Genbank	1222629
Sequence 8637	L40933	Genbank	1160964
Sequence 8638	L41607	Genbank	886293
Sequence 8639	L41816	Genbank	790789
Sequence 8640	L42379	Genbank	1203964
Sequence 8641	L42542	Genbank	974142
Sequence 8642	L42809	Genbank	1160966
Sequence 8643	L43509	Genbank	896282
Sequence 8644	L47162	Genbank	1082035
Sequence 8645	L47647	Genbank	1000861
Sequence 8646	L49345	Genbank	1405422
Sequence 8647	L54057	Genbank	1196416
Sequence 8648	L77701	Genbank	1280205
Sequence 8649	L78464	Genbank	1479980
Sequence 8650	L78669	Genbank	1374862
Sequence 8651	M11147	Genbank	182513
Sequence 8652	M11560	Genbank	178350
Sequence 8653	M12938	Genbank	182515
Sequence 8654	M13520	Genbank	/ <b>1794</b> 59
Sequence 8655	M13755	Genbank	184570
Sequence 8656	M14326	Genbank	340005
Sequence 8657	M14631	Genbank	183416
Sequence 8658	M15182	Genbank	183232
Sequence 8659	M16038	Genbank	187268
Sequence 8660	M16247	Genbank	178044
Sequence 8661	M16279	Genbank	188542
Sequence 8662	M17081	Genbank	178151
Sequence 8663	M17851	Genbank	182860
Sequence 8664	M17885	Genbank	190231
Sequence 8665	M17886	Genbank	190233
Sequence 8666	M18112	Genbank	190166
Sequence 8667	M18157	Genbank	186640
Sequence 8668	M18366	Genbank	179131
Sequence 8669	M19961	Genbank	180940
Sequence 8670	M20372	Genbank	189372
Sequence 8671	M20776	Genbank	180909
Sequence 8672	M21154	Genbank	178517
Sequence 8673	M21575	Genbank	1311702
Sequence 8674	M21895	Genbank	189523
Sequence 8675	M22382	Genbank	190126
Sequence 8676	M22538	Genbank	986883
Sequence 8677	M22803	Genbank	190379
Sequence 8678	M22919	Genbank	189016
Sequence 8679	M22920	Genbank	189021
Sequence 8680	M22960	Genbank	190282
Sequence 8681	M23254	Genbank	511636
Sequence 8682	M23419	Genbank	181996
Sequence 8683	M23725	Genbank	189997
Sequence 8684	M24194	Genbank	187701
Sequence 8685	M24355	Genbank	182604
Sequence 8686	M24486	Genbank	190785

Sequence 8687         M24543         Genbank         341200           Sequence 8688         M26066         Genbank         189854           Sequence 8690         M26663         Genbank         186688           Sequence 8691         M29366         Genbank         181979           Sequence 8692         M29551         Genbank         181979           Sequence 8693         M31159         Genbank         189708           Sequence 8694         M31511         Genbank         179907           Sequence 8696         M31606         Genbank         18940           Sequence 8696         M34055         Genbank         190791           Sequence 8698         M34064         Genbank         190791           Sequence 8699         M34081         Genbank         190791           Sequence 8700         M34088         Genbank         190791           Sequence 8701         M34688         Genbank         190733           Sequence 8701         M34688         Genbank         190738           Sequence 8704         M55153         Genbank         189378           Sequence 8704         M55613         Genbank         189378           Sequence 8706         M57667         G					
Sequence 8688         M26066         Genbank         189854           Sequence 8699         M26325         Genbank         186688           Sequence 8691         M29366         Genbank         618463           Sequence 8692         M29551         Genbank         180708           Sequence 8694         M31159         Genbank         189708           Sequence 8695         M31606         Genbank         189940           Sequence 8696         M33008         Genbank         190791           Sequence 8696         M33008         Genbank         190791           Sequence 8697         M34055         Genbank         190791           Sequence 8698         M34081         Genbank         190791           Sequence 8700         M34088         Genbank         190738           Sequence 8701         M34688         Genbank         182122           Sequence 8702         M37499         Genbank         1807378           Sequence 8704         M55153         Genbank         183378           Sequence 8705         M55618         Genbank         184483           Sequence 8706         M57567         Genbank         184483           Sequence 8708         M59371 <td< td=""><td>Sequence 86</td><td>687</td><td>M24543</td><td>Genbank</td><td>341200</td></td<>	Sequence 86	687	M24543	Genbank	341200
Sequence 8689         M26325         Genbank         186688           Sequence 8690         M26663         Genbank         618463           Sequence 8691         M29366         Genbank         181979           Sequence 8693         M31159         Genbank         183115           Sequence 8694         M31511         Genbank         189940           Sequence 8696         M3308         Genbank         189940           Sequence 8696         M3308         Genbank         416292           Sequence 8696         M34055         Genbank         416292           Sequence 8698         M34081         Genbank         190791           Sequence 8699         M34081         Genbank         190791           Sequence 8700         M34088         Genbank         182122           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34688         Genbank         189378           Sequence 8702         M37499         Genbank         189378           Sequence 8703         M38188         Genbank         189378           Sequence 8704         M55153         Genbank         184483           Sequence 8705         M55618         Ge			M26066	Genbank	189854
Sequence 8690         M26663         Genbank         181979           Sequence 8691         M29366         Genbank         181979           Sequence 8692         M29551         Genbank         180708           Sequence 8693         M31159         Genbank         183115           Sequence 8694         M31511         Genbank         179907           Sequence 8696         M31606         Genbank         190791           Sequence 8698         M34055         Genbank         190791           Sequence 8699         M34081         Genbank         190791           Sequence 8700         M34088         Genbank         190738           Sequence 8700         M34088         Genbank         190738           Sequence 8701         M34688         Genbank         190738           Sequence 8702         M37499         Genbank         189378           Sequence 8704         M55153         Genbank         188378           Sequence 8704         M55153         Genbank         184483           Sequence 8706         M57567         Genbank         184483           Sequence 8706         M57567         Genbank         182591           Sequence 8709         M59349					
Sequence 8691         M29366         Genbank         181979           Sequence 8692         M29551         Genbank         180708           Sequence 8693         M31159         Genbank         183115           Sequence 8694         M31511         Genbank         179907           Sequence 8695         M31606         Genbank         189940           Sequence 8696         M33008         Genbank         190791           Sequence 8697         M34065         Genbank         190791           Sequence 8698         M34081         Genbank         190791           Sequence 8700         M34081         Genbank         190738           Sequence 8700         M34088         Genbank         190738           Sequence 8701         M34688         Genbank         180773           Sequence 8703         M38188         Genbank         189378           Sequence 8703         M38188         Genbank         189378           Sequence 8706         M556153         Genbank         18483           Sequence 8706         M57567         Genbank         184483           Sequence 8707         M58028         Genbank         181943           Sequence 8710         M61715					
Sequence 8692         M29551         Genbank         180708           Sequence 8693         M31159         Genbank         179907           Sequence 8695         M31606         Genbank         189940           Sequence 8696         M33308         Genbank         340236           Sequence 8697         M34055         Genbank         190791           Sequence 8699         M34081         Genbank         13468           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34668         Genbank         182122           Sequence 8703         M34818         Genbank         189378           Sequence 8704         M55153         Genbank         189378           Sequence 8704         M55153         Genbank         189378           Sequence 8706         M55618         Genbank         184483           Sequence 8707         M58028         Genbank         184483           Sequence 8708         M59371         Genbank         181943           Sequence 8708         M59371         Genbank         182591           Sequence 8710         M61715         Genbank         183663           Sequence 8711         M62810         G					
Sequence 8693         M31159         Genbank         179907           Sequence 8694         M31511         Genbank         179907           Sequence 8695         M31606         Genbank         189940           Sequence 8697         M34055         Genbank         340236           Sequence 8698         M34064         Genbank         190791           Sequence 8698         M34064         Genbank         190791           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         189378           Sequence 8705         M55153         Genbank         189378           Sequence 8705         M55618         Genbank         184483           Sequence 8705         M55618         Genbank         184483           Sequence 8706         M57567         Genbank         184483           Sequence 8707         M58028         Genbank         184483           Sequence 8709         M59371         Genbank         182591           Sequence 8711         M62810         Genbank         183861           Sequence 8711         M62810					
Sequence 8694         M31511         Genbank         178907           Sequence 8695         M31606         Genbank         189940           Sequence 8697         M34055         Genbank         340236           Sequence 8698         M34064         Genbank         190791           Sequence 8700         M34081         Genbank         337458           Sequence 8700         M34081         Genbank         190738           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         189378           Sequence 8703         M38188         Genbank         189378           Sequence 8703         M38188         Genbank         189378           Sequence 8706         M55153         Genbank         18483           Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         184483           Sequence 8708         M59371         Genbank         182591           Sequence 8709         M58402         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         G					
Sequence 8695         M31606         Genbank         340236           Sequence 8696         M33008         Genbank         340236           Sequence 8697         M34055         Genbank         190791           Sequence 8698         M34081         Genbank         416292           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34668         Genbank         189378           Sequence 8702         M37499         Genbank         189378           Sequence 8703         M38188         Genbank         189378           Sequence 8704         M55153         Genbank         189378           Sequence 8705         M55618         Genbank         14483           Sequence 8706         M55618         Genbank         148483           Sequence 8707         M58028         Genbank         148483           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8711         M62810         Genbank         182591           Sequence 8712         M64098         Genbank         183891           Sequence 8714         M64788         G					
Sequence 8696         M33308         Genbank         340236           Sequence 8698         M34064         Genbank         190791           Sequence 8699         M34081         Genbank         337458           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         189378           Sequence 8702         M37499         Genbank         189378           Sequence 8702         M37499         Genbank         189378           Sequence 8704         M55153         Genbank         189378           Sequence 8706         M55618         Genbank         184483           Sequence 8706         M57567         Genbank         184483           Sequence 8706         M559371         Genbank         340071           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         183891           Sequence 8713         M64571         Genbank         183891           Sequence 8714         M64780 <td< td=""><td></td><td></td><td></td><td></td><td></td></td<>					
Sequence 8697         M34055         Genbank         190791           Sequence 8698         M34064         Genbank         416292           Sequence 8700         M34081         Genbank         182122           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         188886           Sequence 8703         M38188         Genbank         189378           Sequence 8704         M55153         Genbank         189378           Sequence 8706         M55618         Genbank         184483           Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         181943           Sequence 8708         M59371         Genbank         182591           Sequence 8708         M59371         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         183663           Sequence 8711         M62810         Genbank         183891           Sequence 8713         M64571         Genbank         183891           Sequence 8714         M64788					
Sequence 8698         M34064         Genbank         416292           Sequence 8699         M34081         Genbank         337468           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         188886           Sequence 8704         M55153         Genbank         189378           Sequence 8705         M55618         Genbank         178986           Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         184483           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         183891           Sequence 8711         M62810         Genbank         183891           Sequence 8713         M64571         Genbank         183891           Sequence 8714         M64788         Genbank         190855           Sequence 8716         M65217         Genbank         190364           Sequence 8718         M69181					
Sequence 8699         M34081         Genbank         337458           Sequence 8700         M34088         Genbank         182122           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         188886           Sequence 8704         M55153         Genbank         339520           Sequence 8706         M55618         Genbank         184483           Sequence 8706         M55667         Genbank         184483           Sequence 8707         M58028         Genbank         340071           Sequence 8709         M59849         Genbank         181943           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         183891           Sequence 8712         M64098         Genbank         183891           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         190855           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         190364           Sequence 8718         M69181					
Sequence 8700         M34088         Genbank         192122           Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         188886           Sequence 8703         M38188         Genbank         189378           Sequence 8704         M55153         Genbank         339520           Sequence 8706         M57567         Genbank         184483           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         340071           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         182591           Sequence 8711         M62810         Genbank         183891           Sequence 8711         M64098         Genbank         183891           Sequence 8714         M64788         Genbank         190855           Sequence 8714         M64780         Genbank         190364           Sequence 8719         M73816         Genbank         190364           Sequence 8720         M74491					
Sequence 8701         M34668         Genbank         190738           Sequence 8702         M37499         Genbank         188886           Sequence 8703         M38188         Genbank         189378           Sequence 8704         M55153         Genbank         339520           Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         183681           Sequence 8711         M62810         Genbank         183881           Sequence 8711         M64098         Genbank         183881           Sequence 8714         M64788         Genbank         190855           Sequence 8714         M64788         Genbank         190364           Sequence 8716         M67480         Genbank         184044           Sequence 8719         M73816         Genbank         14957           Sequence 8720         M74491         G					
Sequence 8702         M37499         Genbank         188886           Sequence 8704         M38188         Genbank         189378           Sequence 8705         M55618         Genbank         184483           Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8710         M61715         Genbank         182591           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190865           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491					
Sequence 8704         M38188         Genbank         339520           Sequence 8705         M55618         Genbank         339520           Sequence 8706         M5567         Genbank         184483           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8712         M64098         Genbank         183891           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         190855           Sequence 8716         M67480         Genbank         190364           Sequence 8719         M73816         Genbank         184025           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M76491         Genbank         184020           Sequence 8721         M76729         G					
Sequence 8704         M55153         Genbank         339520           Sequence 8705         M55618         Genbank         184483           Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190365           Sequence 8716         M67480         Genbank         190364           Sequence 8716         M69181         Genbank         184404           Sequence 8719         M73816         Genbank         183609           Sequence 8719         M74491         Genbank         184020           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M76729					
Sequence 8706         M57567         Genbank         184483           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         183663           Sequence 8711         M62810         Genbank         183891           Sequence 8712         M64098         Genbank         183891           Sequence 8714         M64788         Genbank         187382           Sequence 8715         M65217         Genbank         190855           Sequence 8716         M65217         Genbank         190364           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         183609           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M75126         Genbank         181273           Sequence 8723         M80254	Sequence 8			Genbank	
Sequence 8706         M57567         Genbank         178986           Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         188563           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         190855           Sequence 8714         M64788         Genbank         190855           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         188625           Sequence 8718         M69181         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8721         M75126         Genbank         184020           Sequence 8723         M80254         Genbank         189519           Sequence 8724         M80335			M55153		
Sequence 8707         M58028         Genbank         340071           Sequence 8708         M59371         Genbank         181943           Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         190855           Sequence 8714         M64788         Genbank         190855           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         190364           Sequence 8718         M69181         Genbank         188625           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M75126         Genbank         184020           Sequence 8723         M80254         Genbank         189519           Sequence 8724         M80335         Genbank         18072           Sequence 8725         M80646         G			M55618	Genbank	184483
Sequence 8708         M59371         Genbank         181943           Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         190364           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         183609           Sequence 8718         M69181         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M76729         Genbank         184020           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         181273           Sequence 8725         M80646	Sequence 8	706	M57567	Genbank	178986
Sequence 8708         M59371         Genbank         181943           Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         190364           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         183609           Sequence 8718         M69181         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M76729         Genbank         184020           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         181273           Sequence 8725         M80646			M58028	Genbank	340071
Sequence 8709         M59849         Genbank         182591           Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         190364           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         183609           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M75126         Genbank         189519           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         18956           Sequence 8725         M80646         Genbank         18976           Sequence 8726         M80776         Ge			M59371	Genbank	181943
Sequence 8710         M61715         Genbank         340367           Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         184404           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         190364           Sequence 8718         M69181         Genbank         188625           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8720         M74491         Genbank         184020           Sequence 8721         M75126         Genbank         184020           Sequence 8723         M80254         Genbank         181273           Sequence 8723         M80254         Genbank         181273           Sequence 8726         M80776			M59849	Genbank	182591
Sequence 8711         M62810         Genbank         188563           Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         184404           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         184205           Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M76729         Genbank         184020           Sequence 8723         M80254         Genbank         189519           Sequence 8724         M80335         Genbank         189519           Sequence 8724         M80335         Genbank         189519           Sequence 8725         M80646         Genbank         18071           Sequence 8726         M80776         G			M61715	Genbank	340367
Sequence 8712         M64098         Genbank         183891           Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         184404           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         190364           Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M75126         Genbank         189519           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80335         Genbank         181273           Sequence 8724         M80335         Genbank         18966           Sequence 8725         M80646         Genbank         17934           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Ge			M62810	Genbank	188563
Sequence 8713         M64571         Genbank         187382           Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         184404           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M75126         Genbank         189519           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         18071           Sequence 8726         M80776         Genbank         180172           Sequence 8728         M83651         Genbank         180132           Sequence 8730         M84739         G			M64098	Genbank	183891
Sequence 8714         M64788         Genbank         190855           Sequence 8715         M65217         Genbank         184404           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M75126         Genbank         189519           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         189519           Sequence 8723         M80335         Genbank         189519           Sequence 8724         M80335         Genbank         18966           Sequence 8725         M80646         Genbank         18966           Sequence 8725         M80776         Genbank         180172           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Ge				Genbank	
Sequence 8715         M65217         Genbank         184404           Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         183609           Sequence 8721         M75126         Genbank         184020           Sequence 8721         M75126         Genbank         189519           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         189519           Sequence 8723         M80335         Genbank         189519           Sequence 8724         M80335         Genbank         18956           Sequence 8725         M80646         Genbank         189966           Sequence 8725         M80776         Genbank         180172           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         178163           Sequence 8730         M84739         G					
Sequence 8716         M67480         Genbank         190364           Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         641957           Sequence 8720         M74491         Genbank         183609           Sequence 8720         M74491         Genbank         178161           Sequence 8721         M75126         Genbank         184020           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         189966           Sequence 8726         M80776         Genbank         179334           Sequence 8726         M80776         Genbank         180172           Sequence 8728         M83651         Genbank         180172           Sequence 8729         M84326         Genbank         178163           Sequence 8731         M85038         Genbank         179881           Sequence 8731         M85038         Genbank         180189           Sequence 8733         M91670					
Sequence 8717         M69066         Genbank         188625           Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         178161           Sequence 8721         M75126         Genbank         184020           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         189966           Sequence 8726         M80776         Genbank         179334           Sequence 8726         M80493         Genbank         180172           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         180195           Sequence 8734         M95627				• • • • • • • • • • • • • • • • • • • •	
Sequence 8718         M69181         Genbank         641957           Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         178161           Sequence 8721         M75126         Genbank         184020           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         189519           Sequence 8723         M80254         Genbank         189519           Sequence 8724         M80335         Genbank         189519           Sequence 8724         M80335         Genbank         189566           Sequence 8725         M80646         Genbank         18966           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         178163           Sequence 8729         M84326         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8731         M85038         Genbank         180189           Sequence 8733         M91670         G					
Sequence 8719         M73816         Genbank         183609           Sequence 8720         M74491         Genbank         178161           Sequence 8721         M75126         Genbank         184020           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         180172           Sequence 8729         M84326         Genbank         179881           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         180749           Sequence 8736         M96803         Genbank         190749           Sequence 8738         M97501	•			•	
Sequence 8720         M74491         Genbank         178161           Sequence 8721         M75126         Genbank         184020           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         13032           Sequence 8729         M84326         Genbank         179881           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8733         M91670         Genbank         183369           Sequence 8734         M95627         Genbank         181915           Sequence 8736         M96803         Genbank         190749           Sequence 8737         M96995         Genbank         180621           Sequence 8739         M97934         G					
Sequence 8721         M75126         Genbank         184020           Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         180172           Sequence 8729         M84326         Genbank         179881           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         190749           Sequence 8736         M96803         Genbank         190749           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501					
Sequence 8722         M76729         Genbank         189519           Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         431032           Sequence 8729         M84326         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         180189           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         190749           Sequence 8736         M96803         Genbank         180749           Sequence 8737         M96995         Genbank         180621           Sequence 8738         M97501         Genbank         180621           Sequence 8740         M97935					
Sequence 8723         M80254         Genbank         181273           Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         431032           Sequence 8729         M84326         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         190749           Sequence 8735         M96684         Genbank         190749           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935					
Sequence 8724         M80335         Genbank         189966           Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         431032           Sequence 8729         M84326         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         190749           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478 <td< td=""><td></td><td></td><td></td><td></td><td></td></td<>					
Sequence 8725         M80646         Genbank         338701           Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         431032           Sequence 8729         M84326         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         180621           Sequence 8738         M97501         Genbank         180621           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8726         M80776         Genbank         179334           Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         431032           Sequence 8729         M84326         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         180621           Sequence 8738         M97501         Genbank         180621           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577				<b>—</b>	
Sequence 8727         M81934         Genbank         180172           Sequence 8728         M83651         Genbank         431032           Sequence 8729         M84326         Genbank         178163           Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         180621           Sequence 8738         M97501         Genbank         180621           Sequence 8740         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence         8728         M83651         Genbank         431032           Sequence         8729         M84326         Genbank         178163           Sequence         8730         M84739         Genbank         179881           Sequence         8731         M85038         Genbank         190189           Sequence         8732         M86492         Genbank         183369           Sequence         8733         M91670         Genbank         181915           Sequence         8734         M95627         Genbank         870802           Sequence         8735         M96684         Genbank         190749           Sequence         8736         M96803         Genbank         338442           Sequence         8737         M96995         Genbank         180621           Sequence         8738         M97501         Genbank         180621           Sequence         8739         M97934         Genbank         186566           Sequence         8740         M97935         Genbank         2281070           Sequence         8741         M98478         Genbank         339577					
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Sequence 8730         M84739         Genbank         179881           Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8731         M85038         Genbank         190189           Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8732         M86492         Genbank         183369           Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					,
Sequence 8733         M91670         Genbank         181915           Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577	Sequence 8	/31			
Sequence 8734         M95627         Genbank         870802           Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8735         M96684         Genbank         190749           Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8736         M96803         Genbank         338442           Sequence 8737         M96995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577			::::::::::::::::::::::::::::::::::::::		
Sequence 8737         M98995         Genbank         181975           Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8738         M97501         Genbank         180621           Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8739         M97934         Genbank         186566           Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577					
Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577			M97501	Genbank	
Sequence 8740         M97935         Genbank         2281070           Sequence 8741         M98478         Genbank         339577	Sequence 8	739	M97934	Genbank	186566
Sequence 8741 M98478 Genbank 339577			M97935	Genbank :	2281070
			M98478	Genbank	339577
			M99422	Genbank	183662

Table 3-3

Sequence 8743	S60710	Genbank	300301
Sequence 8744	S62138	Genbank	38 <b>6158</b>
Sequence 8745	S67779	Genbank	459372
Sequence 8746	S69272	Genbank	546087
Sequence 8747	S70114	Genbank	546696
Sequence 8748	S73498	Genbank	688010
Sequence 8749	S74678	Genbank	241477
Sequence 8750	S74799	Genbank	797365
Sequence 8751	S75174	Genbank	802120
Sequence 8752	S82081	Genbank	1488412
Sequence 8753	U00930	Genbank	
Sequence 8754	U00947	Genbank	405043
Sequence 8755	U02390		405049
Sequence 8756	U02390	Genbank Genbank	409928
			407307
Sequence 8757	U03886	Genbank	458225
Sequence 8758	U05572	Genbank	1419373
Sequence 8759	U07563	Genbank	514264
Sequence 8760	U07681	Genbank	706838
Sequence 8761	U07810	Genbank	469038
Sequence 8762	U07919	Genbank	995 <b>897</b>
Sequence 8763	U08377	Genbank	50 <b>8230</b>
Sequence 8764	U09220	Genbank	79 <b>9338</b>
Sequence 8765	U09284	Genbank	51 <b>6011</b>
Sequence 8766	U09716	Genbank	60 <b>6827</b>
Sequence 8767	U09813	Genbank	1008454
Sequence 8768	U10248	Genbank	984 <b>280</b>
Sequence 8769	U10323	Genbank	532 <b>312</b>
Sequence 8770	U10439	Genbank	5771 <b>69</b>
Sequence 8771	U10691	Genbank	533 <b>522</b>
Sequence 8772	U10886	Genbank	1685074
Sequence 8773	U12424	Genbank	52 <b>5319</b>
Sequence 8774	U12535	Genbank	530822
Sequence 8775	U13022	Genbank	537 <b>293</b>
Sequence 8776	U13991	Genbank	562 <b>076</b>
Sequence 8777	U14193	Genbank	5 <b>55899</b>
Sequence 8778	U14966	Genbank	55 <b>0012</b>
Sequence 8779	U14969	Genbank	550018
Sequence 8780	U14970	Genbank	5500 <b>20</b>
Sequence 8781	U14971	Genbank	550 <b>022</b>
Sequence 8782	U15158	Genbank	576 <b>622</b>
Sequence 8783	U16660	Genbank	56 <b>4064</b>
Sequence 8784	U18121	Genbank	915 <b>283</b>
Sequence 8785	U18197	Genbank	603 <b>073</b>
Sequence 8786	U18937	Genbank	89 <b>9108</b>
Sequence 8787	U19145	Genbank	914904
Sequence 8788	U19252	Genbank	80 <b>8035</b>
Sequence 8789	U20476	Genbank	3077819
Sequence 8790	U21910	Genbank	736 <b>402</b>
Sequence 8791	U22055	Genbank	79 <b>9176</b>
Sequence 8792	U22815	Genbank	930340
Sequence 8793	U23028	Genbank	806853
Sequence 8794	U24169	Genbank	1215668
Sequence 8795	U24223	Genbank	1215670
Sequence 8796	U25816	Genbank	837262
Sequence 8797	U26425	Genbank	836664
Sequence 8798	U27515	Genbank	4096274
			•

Sequence 8799	U28387	Genbank	881950
Sequence 8800	U28686	Genbank	881953
Sequence 8801	U28727	Genbank	1142969
Sequence 8802	U28811	Genbank	1373018
Sequence 8803	U30255	Genbank	984324
Sequence 8804	U32944	Genbank	1209060
Sequence 8805	U32989	Genbank	993045
Sequence 8806	U33460	Genbank	4096590
Sequence 8807	U33760	Genbank	995823
Sequence 8808	U33818	Genbank	1163176
Sequence 8809	U34038	Genbank	1041728
Sequence 8810	U34877	Genbank	1143231
Sequence 8811	U34994	Genbank	6525311
Sequence 8812	U35116	Genbank	1008541
Sequence 8813	U36188	Genbank	1244507
Sequence 8814	U36764	Genbank	1036804
	U37426	Genbank	1171152
Sequence 8815		Genbank	
Sequence 8816	U37558	• • • • • • • • • • • • • • • • • • • •	1017812
Sequence 8817	U38784	Genbank	1574947
Sequence 8818	U38894	Genbank	1589739
Sequence 8819	U38945	Genbank	1353569
Sequence 8820	U39400	Genbank	1234796
Sequence 8821	U39945	Genbank	1209686
Sequence 8822	U40152	Genbank	1113100
Sequence 8823	U40989	Genbank	1125005
Sequence 8824	U41371	Genbank	1173904
Sequence 8825	U41387	Genbank	1230563
Sequence 8826	U41806	Genbank	1145798
Sequence 8827	U46691	Genbank (	1401077
Sequence 8828	U47742	Genbank	1517913
Sequence 8829	U49436	Genbank	1229139
Sequence 8830	U50079	Genbank	1277083
Sequence 8831	U51587	Genbank	1669823
Sequence 8832	U51625	Genbank	1695798
Sequence 8833	U51678	Genbank	1915966
Sequence 8834	U53209	Genbank	1256836
Sequence 8835	U53346	Genbank	1545981
Sequence 8836	U53530	Genbank	1314642
Sequence 8837	U54559	Genbank	2351379
Sequence 8838	U57052	Genbank	1519039
Sequence 8839	U58766	Genbank	1381178
Sequence 8840	U59435	Genbank	2697004
Sequence 8841	U61262	Genbank	1621606
Sequence 8842	U61397	Genbank	1518693
Sequence 8843	U61734	Genbank	1407825
Sequence 8844	U61843	Genbank	3650451
Sequence 8845	U62317	Genbank	6862558
Sequence 8846	U62583	Genbank	1778050
Sequence 8847	U63296	Genbank	2047312
Sequence 8848	U63743	Genbank	1695881
Sequence 8849	U64791	Genbank	1498734
Sequence 8850	U65676	Genbank	1654350
Sequence 8851	U65928	Genbank	1549382
Sequence 8852	U66618	Genbank	1549362
Sequence 8853	U67058	Genbank	4097702
	U68723	Genbank	2114391
Sequence 8854	000123	Genbank	2114381

Table 3-3

Sequence 8855	U700 <b>63</b>	Genbank	1743866
Sequence 8856	U70312	Genbank	2865218
Sequence 8857	U70451	Genbank	1763090
Sequence 8858	U72069	Genbank	1657775
Sequence 8859	U72398	Genbank	1622940
Sequence 8860	U72515	Genbank	1673519
Sequence 8861	U73167	Genbank	2880032
Sequence 8862	U73338	Genbank	1763268
Sequence 8863	U74612	Genbank	1842252
Sequence 8864	U74613	Genbank	1842254
Sequence 8865	U75329	Genbank	2507612
Sequence 8866	U75330	Genbank	2507614
Sequence 8867	U76638	Genbank	1710174
Sequence 8868	U76764	Genbank	1685050
Sequence 8869	U77456	Genbank	1679778
Sequence 8870	U79241	Genbank	1710187
Sequence 8871	U79302	Genbank	1710287
Sequence 8872	U79458	Genbank	4205085
Sequence 8873	U80040	Genbank	1718501
Sequence 8874	U80735	Genbank	2565045
Sequence 8875	U80811	Genbank	1857424
Sequence 8876	U81002	Genbank	4580010
Sequence 8877	U81599	Genbank	1764089
Sequence 8878	U82226	Genbank	1848270
Sequence 8879	U82535	Genbank	2149155
Sequence 8880	U83410	Genbank	1923242
Sequence 8881	U85658	Genbank	2058552
Sequence 8882	U89278	Genbank	1877500
Sequence 8883	U89505	Genbank	2078528
Sequence 8884	U89 <b>995</b>	Genbank	2078532
Sequence 8885	U90339	Genbank	1906010
Sequence 8886	U94747	Genbank	2290529
Sequence 8887	U94788	Genbank	3041866
Sequence 8888	U968 <b>76</b>	Genbank	2358268
Sequence 8889	U97156	Genbank	2459796
Sequence 8890	U97519	Genbank	2213812
Sequence 8891	V00572	Genbank	35434
Sequence 8892	X01060	Genbank	37432
Sequence 8893	X02152	Genbank	34312
Sequence 8894	X03363	Genbank	31197
Sequence 8895	X03473	Genbank	32106
Sequence 8896	X04408	Genbank	31914
Sequence 8897	X04526	Genbank	31667
Sequence 8898	X04741	Genbank	35439
Sequence 8899	X05332	Genbank	35740
Sequence 8900	X06272	Genbank	30865
Sequence 8901	X06409	Genbank	30204
Sequence 8902	X06562	Genbank	31737
Sequence 8903	X06747	Genbank	36101
Sequence 8904	X06994	Genbank	30302
Sequence 8905	X12433	Genbank	32451
Sequence 8906	X12597	Genbank	32326
Sequence 8907	X12660	Genbank	36654
Sequence 8908	X13294	Genbank	28685
Sequence 8909	X13916	Genbank	34338
Sequence 8910	X139 <b>39</b>	Genbank	34237

Table 3-3

Sequence 8911	X15488	Genbank	30933
Sequence 8912	X15729	Genbank	38317
Sequence 8913	X15880	Genbank	30029
Sequence 8914	X16064	Genbank	37495
Sequence 8915	X16135	Genbank	32355
Sequence 8916	X16396	Genbank	35070
Sequence 8917	X17567	Genbank	36512
Sequence 8918	X17620	Genbank	35067
Sequence 8919	X51466	Genbank	31105
Sequence 8920	X52519	Genbank	36694
Sequence 8921	X52851	Genbank	30167
Sequence 8922	X52955	Genbank	29627
Sequence 8923	X53280	Genbank	29504
Sequence 8924	X56998	Genbank	37564
Sequence 8925	X57129	Genbank	31967
Sequence 8926	X57206	Genbank	33990
Sequence 8927	X59618	Genbank	36154
Sequence 8928	X61123	Genbank	29508
Sequence 8929	X64123	Genbank	35819
Sequence 8930	X64229	Genbank	30502
Sequence 8931	X64707	Genbank	29382
Sequence 8932	X64875	Genbank	398163
Sequence 8933	X66364	Genbank	36620
Sequence 8934	X67155	Genbank	6723674
Sequence 8935	X67334	Genbank	468703
Sequence 8936	X68836	Genbank	36326
Sequence 8937	X70940	Genbank	38455
Sequence 8938	X71428	Genbank	393415
Sequence 8939	X71810	Genbank	297905
Sequence 8940	X74070	Genbank	395086
Sequence 8941	X74794	Genbank	683749
Sequence 8942	X76057	Genbank	416016
Sequence 8943	X78136	Genbank	460772
Sequence 8944	X78706	Genbank	471311
Sequence 8945	X80197	Genbank	951271
Sequence 8946	X80230	Genbank	599828
Sequence 8947	X80910	Genbank	531475
Sequence 8948	X81197	Genbank	773572
Sequence 8949	X81889	Genbank	1702923
Sequence 8950	X82321	Genbank	1617117
Sequence 8951	X85137	Genbank	1155083
Sequence 8952	X86012	Genbank	976082
Sequence 8953	X92814	Genbank	1054751
Sequence 8954	X94333	Genbank	1518268
Sequence 8955	X94754	Genbank	1702931
Sequence 8956	X95762	Genbank	2584786
Sequence 8957	X96698	Genbank	1418766
Sequence 8958	X97064	Genbank	1296663
Sequence 8959	X97124	Genbank	1518264
Sequence 8960	X97335	Genbank	1507823
Sequence 8961	X97674	Genbank	1877214
Sequence 8962	X97795	Genbank	1495482
Sequence 8963	X98054	Genbank	1359754
Sequence 8964	X98261	Genbank	1770459
Sequence 8965	X98507	Genbank	1926310
Sequence 8966	X99141	Genbank	1903215
20422.000000			

Table 3-3

Sequence 8967	X99802	Genbank	2769561
Sequence 8968	Y00052	Genbank	30308
Sequence 8969	Y00281	Genbank	36052
Sequence 8970	Y00282	Genbank	36048
Sequence 8971	Y00815	Genbank	34266
Sequence 8972	Y00978	Genbank	35359
Sequence 8973	Y08134	Genbank	1552274
Sequence 8974	Y09862	Genbank	1945760
Sequence 8975	Y10275	Genbank	1890330
Sequence 8976	Y12478	Genbank	1946204
Sequence 8977	Y13621	Genbank	2190399
Sequence 8978	Y13936	Genbank	2315201
Sequence 8979	Y14486	Genbank	5830437
Sequence 8980	Y15286	Genbank	2584788
Sequence 8981	Y18004	Genbank	4490941
Sequence 8982	Z11583	Genbank	35118
Sequence 8983	Z14244	Genbank	30150
Sequence 8984	Z18956	Genbank	36726
Sequence 8985	Z26649	Genbank	510116
Sequence 8986	Z37166	Genbank	587145
Sequence 8987	Z46376	Genbank	587201
Sequence 8988	Z50022	Genbank	1107702
Sequence 8989	Z58902	Genbank	1030815
Sequence 8990	Z69709	Genbank	1204107
Sequence 8991	Z69780	Genbank	1514560
	Z70200	Genbank	3255964
Sequence 8992		Genbank	6580491
Sequence 8993	Z71183	Genbank	1545951
Sequence 8994	Z72499		1322149
Sequence 8995	Z73358	Genbank	
Sequence 8996	Z73639	Genbank	2632099
Sequence 8997	Z83840	Genbank	4914518
Sequence 8998	Z84466	Genbank	3319673
Sequence 8999	Z92543	Genbank	3164068
Sequence 9000	Z92545	Genbank	2462414
Sequence 9001	Z93783	Genbank	4581336
Sequence 9002	Z95114	Genbank	5101742
Sequence 9003	Z97053	Genbank	9650676
Sequence 9004	Z98258	Genbank	2462407
Sequence 9005	Z98752	Genbank	9408718
Sequence 9006	Z99716	Genbank	4456457
Sequence 9007	AA011002	EST	1472029
Sequence 9008	AA024935	EST	1489859
Sequence 9009	AA025057	EST	1489962
Sequence 9010	AA025381	EST	1491520
Sequence 9011	AA027341	EST	1492958
Sequence 9012	AA031969	EST	1501922
Sequence 9013	AA033521	EST	1505367
Sequence 9014	AA033724	EST	1505542
Sequence 9015	AA037091	EST	1512209
Sequence 9016	AA037126	EST	1512234
Sequence 9017	AA037364	EST	1512602
Sequence 9018	AA040182	EST	1516598
Sequence 9019	AA043058	EST	1522593
Sequence 9020	AA044122	EST	1521980
Sequence 9021	AA044586	EST	1522977
Sequence 9022	AA044813	EST	1523016
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Table 3-3

Sequence 9023	AA047046	EST	1524944
Sequence 9024	AA053156	EST	1544427
Sequence 9025	AA053431	EST	1544086
Sequence 9026	AA053679	EST	1544606
Sequence 9027	AA054463	EST	1545388
Sequence 9028	AA055407	EST	1547946
Sequence 9029	AA055649	EST	1547988
Sequence 9030	AA056099	EST	
Sequence 9031	AA056273		1548436
Sequence 9032	AA056358	EST EST	1548820
Sequence 9033	AA058929		1548698
		EST	1551784
Sequence 9034	AA062553	EST	1556727
Sequence 9035	AA065298	EST .	1929298
Sequence 9036	AA069599	EST	1576958
Sequence 9037	AA070333	EST	1577692
Sequence 9038	AA071122	EST	1578482
Sequence 9039	AA074907	EST	1614776
Sequence 9040	AA075864	EST	1615735
Sequence 9041	AA075873	EST	1615744
Sequence 9042	AA075984	EST	1615853
Sequence 9043	AA077364	EST	1836838
Sequence 9044	AA081041	EST	1622976
Sequence 9045	AA081608	EST	1623666
Sequence 9046	AA082133	EST	1624191
Sequence 9047	AA082281	EST	1624356
Sequence 9048	AA082487	EST	1624560
Sequence 9049	AA082543	EST	1624600
Sequence 9050	AA082817	EST	1624874
Sequence 9051	AA083182	EST	1625258
Sequence 9052	AA083288	EST	1625349
Sequence 9053	AA083300	EST	1625361
Sequence 9054	AA088420	EST	1633967
Sequence 9055	AA098867	EST	1645051
Sequence 9056	AA099415	EST	1645308
Sequence 9057	AA099496	EST	1645449
Sequence 9058	AA099923	EST	1646071
Sequence 9059	AA099926	EST	1646074
Sequence 9060	AA101077	EST	1647597
Sequence 9061	AA101100	EST	1647676
Sequence 9062	AA101277	EST	1648016
Sequence 9063	AA101440	EST	1648254
Sequence 9064	AA111991	EST	1664113
Sequence 9065	AA112480	EST	1665157
Sequence 9066	AA112633	EST	1665334
Sequence 9067	AA112679	EST	1665397
Sequence 9068	AA115409		
Sequence 9069	AA121091	EST	1670591
		EST	1678635
Sequence 9070 Sequence 9071	AA121210 AA126105	EST	1678949
		EST	1685755
Sequence 9072	AA126762	EST	1686262
Sequence 9073	AA128709	EST	1688617
Sequence 9074	AA129754	EST	1690164
Sequence 9075	AA130211	EST	1691215
Sequence 9076	AA130220	EST	1691224
Sequence 9077	AA130247	EST	1691251
Sequence 9078	AA130520	EST	1691803

Table 3-3

Sequence 9079	AA131227	EST	1692735
Sequence 9080	AA131252	EST	1692814
Sequence 9081	AA131474	EST	1692962
Sequence 9082	AA131496	EST	1692984
Sequence 9083	AA132133	EST	1693667
Sequence 9084	AA132163	EST	1693672
Sequence 9085	AA132328	EST	1693818
Sequence 9086	AA132414	EST	1693905
Sequence 9087	AA132885	EST	1694372
Sequence 9088	AA135266	EST	1696351
Sequence 9089	AA135745	EST	1696757
Sequence 9090	AA135830	EST	
Sequence 9091	AA143151	EST	1697090 1712521
Sequence 9092	AA143301	EST	1712672
Sequence 9093	AA147806	EST	1717195
Sequence 9094	AA148139	EST	1717530
Sequence 9095	AA149520	EST	1720338
Sequence 9096	AA149637	EST	1720438
Sequence 9097	AA149750	EST	1720892
Sequence 9098	AA150002	EST	1721155
Sequence 9099	AA150076	EST	1721298
Sequence 9100	AA150383	EST	1721914
Sequence 9101	AA151213	EST	1719468
Sequence 9102	AA151345	EST	1719890
Sequence 9103	AA151506	EST	1720046
Sequence 9104	AA152172	EST	1721224
Sequence 9105	AA156215	EST	1727895
Sequence 9106	AA1568 <b>07</b>	EST	17 <b>284</b> 40
Sequence 9107	AA157 <b>71</b> 5	EST	1729340
Sequence 9108	AA158196	EST	1733007
Sequence 9109	AA158600	EST	1733436
Sequence 9110	AA160097	EST	1734545
Sequence 9111	AA160948	EST	1735663
Sequence 9112	AA164729	EST	1740889
Sequence 9113	AA165403	EST	1741436
Sequence 9114	AA165418	EST	1741554
Sequence 9115	AA167191	EST	1745698
Sequence 9116	AA167829	EST	1744980
Sequence 9117	AA169752	EST	1748137
Sequence 9118	AA171588	EST	1750657
Sequence 9119	AA174204	EST	1754484
Sequence 9120	AA179201	EST	1760816
Sequence 9121	AA179430	EST	1760782
Sequence 9122	AA179442	EST	1760794
Sequence 9123	AA179690	EST	1761923
Sequence 9124	AA181608	EST	1765136
Sequence 9125	AA181618	EST	1765085
Sequence 9126	AA186685	EST	1774906
Sequence 9127	AA186730	EST	1774847
Sequence 9128	AA187424	EST	1773633
Sequence 9129	AA187428	EST	1773747
Sequence 9130	AA187809		
Sequence 9131		EST	1774003
	AA188497	EST	1775584
Sequence 9132 Sequence 9133	AA188499	EST	1775586
•	AA188875	EST	1775902
Sequence 9134	AA19 <b>0915</b>	EST	1779571

Table 3-3

Sequence 9135	AA191160	EST	1779 <b>854</b>
Sequence 9136	AA194169	EST	1783 <b>860</b>
Sequence 9137	AA194896	EST	1784 <b>658</b>
Sequence 9138	AA195259	EST	1784959
Sequence 9139	AA195352	EST	1785043
			1785213
Sequence 9140	AA195510	EST	
Sequence 9141	AA195535	EST	1785248
Sequence 9142	AA196350	EST	1791925
Sequence 9143	AA197018	EST	1792634
Sequence 9144	AA197063	EST	1792 <b>629</b>
Sequence 9145	AA197137	EST	1792 <b>81</b> 1
Sequence 9146	AA203450	EST	1799 <b>16</b> 1
Sequence 9147	AA205846	EST	1801217
Sequence 9148	AA205928	EST	1801316
Sequence 9149	AA206583	EST	1801963
Sequence 9150	AA207229	EST	1802920
Sequence 9151	AA209239	EST	1807254
Sequence 9152	AA210796	EST	1809450
Sequence 9153	AA211864	EST	1810491
Sequence 9154	AA215319	EST	1815082
Sequence 9155	AA215526	EST	1815298
Sequence 9156	AA215534	EST	1815288
Sequence 9157	AA216769	EST	1817460
Sequence 9158	AA218567	EST	1832642
Sequence 9159	AA218697	EST	1832789
Sequence 9160	AA223382	EST	1843907
Sequence 9161	AA225008	EST	1846317
Sequence 9162	AA226737	EST	18483 <b>08</b>
Sequence 9163	AA227227	EST	1848772
Sequence 9164	AA227608	EST	1849 <b>152</b>
Sequence 9165	AA232784	EST	18559 <b>86</b>
Sequence 9166	AA233063	EST	185607 <b>5</b>
Sequence 9167	AA233297	EST	1856 <b>290</b>
Sequence 9168	AA234499	EST	18589 <b>90</b>
Sequence 9169	AA234963	EST	185 <b>8132</b>
Sequence 9170	AA236353	EST	18584 <b>03</b>
Sequence 9171	AA236418	EST	1858 <b>543</b>
Sequence 9172	AA243646	EST	1874456
Sequence 9173	AA243672	EST	18746 <b>55</b>
Sequence 9174	AA247773	EST	18798 <b>78</b>
Sequence 9175	AA252002	EST	18869 <b>63</b>
Sequence 9176	AA252445	EST	1887416
Sequence 9177	AA252465	EST	1887436
Sequence 9178	AA252567	EST	1887541
Sequence 9179	AA252657	EST	1887622
Sequence 9180	AA255982	EST	1891658
Sequence 9181	AA257153	EST	1891281
Sequence 9182	AA258425	EST	1893549
Sequence 9183	AA258503	EST	18936 <b>28</b>
Sequence 9184	AA259061	EST	1894541
Sequence 9185	AA262003	EST	1897984
Sequence 9186	AA262559		
Sequence 9187	AA278264	EST	1898011
•		EST	1920222
Sequence 9188	AA278642	EST	1919962
Sequence 9189	AA278910	EST	1920376
Sequence 9190	AA279735	EST	1921 <b>200</b>

Table 3-3

Sequence 9191	AA280710	EST	1923603
Sequence 9192	AA281238	EST	1923919
Sequence 9193	AA282006	EST	1924848
Sequence 9194	AA282870	EST	1925786
Sequence 9195	AA282920	EST	1925896
Sequence 9196	AA284386	EST	1928720
Sequence 9197	AA284870	EST	1927464
Sequence 9198	AA287235	EST	1934261
Sequence 9199	AA287990	EST	1933814
Sequence 9200	AA291446	EST	1939425
Sequence 9201	AA291926	EST	1940090
Sequence 9202	AA293348	EST	1940745
Sequence 9203	AA294883	EST	1947228
Sequence 9204	AA296484	EST	1948818
Sequence 9205	AA297767	EST	1950170
Sequence 9206	AA299346	EST	1951678
Sequence 9207	AA300574	EST	1953135
Sequence 9208	AA300728	EST	1953296
Sequence 9209	AA301138	EST	1953491
Sequence 9210	AA301439	EST	1953773
Sequence 9211	AA302601	EST	1955083
Sequence 9212	AA303657	EST	1956009
Sequence 9213	AA304899	EST	1957247
Sequence 9214	AA304904	EST	1957469
Sequence 9215	AA304981	EST	1957328
Sequence 9216	AA305193	EST	1957520
Sequence 9217	AA305525	EST	1957850
Sequence 9218	AA30556B	EST	1957915
Sequence 9219	AA305862	EST	1958192
Sequence 9220	AA305873	EST	1958203
Sequence 9221	AA306226	EST	1958554
Sequence 9222	AA306444	EST	1958772
Sequence 9223	AA306511	EST	1958840
Sequence 9224	AA306594	EST	195 <b>8923</b>
Sequence 9225	AA306623	EST	195 <b>8952</b>
Sequence 9226	AA306709	EST	195 <b>9037</b>
Sequence 9227	AA306734	EST	19 <b>59083</b>
Sequence 9228	AA306846	EST	19 <b>59174</b>
Sequence 9229	AA306917	EST	19 <b>59247</b>
Sequence 9230	AA306961	EST	195 <b>9291</b>
Sequence 9231	AA306966	EST	195 <b>9316</b>
Sequence 9232	AA307060	EST	195 <b>9538</b>
Sequence 9233	AA307269	EST	19 <b>59597</b>
Sequence 9234	AA307327	EST	195 <b>9655</b>
Sequence 9235	AA307332	EST	195 <b>9660</b>
Sequence 9236	AA307779	EST	196 <b>0177</b>
Sequence 9237	AA307876	EST	196 <b>0276</b>
Sequence 9238	AA307979	EST	196 <b>0308</b>
Sequence 9239	AA308390	EST	196 <b>0719</b>
Sequence 9240	AA308574	EST	1960922
Sequence 9241	AA309051	EST	196 <b>1377</b>
Sequence 9242	AA309832	EST	19 <b>62325</b>
Sequence 9243	AA310059	EST	196 <b>2388</b>
Sequence 9244	AA310330	EST	196 <b>2679</b>
Sequence 9245	AA311461	EST	19 <b>63787</b>
Sequence 9246	AA311501	EST	196 <b>4046</b>

Sequence 9247	AA311933	EST	1964261
Sequence 9248	AA312489	EST	1964818
Sequence 9249	AA312669	EST	1965018
Sequence 9250	AA312778	EST	1965126
Sequence 9251	AA312886	EST	1965276
Sequence 9252	AA313319	EST	1965648
Sequence 9253	AA313489	EST	1965818
Sequence 9254	AA313896	EST	1966225
Sequence 9255	AA314146	EST	1966495
Sequence 9256	AA314350	EST	1966679
Sequence 9257	AA314550	EST	1966899
Sequence 9258	AA314707	EST	1967056
Sequence 9259	AA314717	EST	1967277
	AA314882	EST	1967452
Sequence 9260	AA315444		1967923
Sequence 9261	AA315472	EST EST	1967801
Sequence 9262		EST	
Sequence 9263	AA315499		1967828
Sequence 9264	AA315975	EST	1968304
Sequence 9265	AA316030	EST	1968359
Sequence 9266	AA316470	EST	1968799
Sequence 9267	AA316551	EST	1968879
Sequence 9268	AA316590	EST	1968918
Sequence 9269	AA316620	EST	1968948
Sequence 9270	AA316641	EST	1968989
Sequence 9271	AA316743	EST	1969142
Sequence 9272	AA316864	EST	1969192
Sequence 9273	AA317246	EST	1969573
Sequence 9274	AA318236	EST	1970564
Sequence 9275	AA318266	EST	1970594
Sequence 9276	AA323846	EST	1976173
Sequence 9277	AA325370	EST	1977838
Sequence 9278	AA325474	EST	1977717
Sequence 9279	AA326465	EST	1978709
Sequence 9280	AA326466	EST	1978710
Sequence 9281	AA328435	EST	1980679
Sequence 9282	AA329233	EST	1981477
Sequence 9283	AA330793	EST	1983035
Sequence 9284	AA332288	EST	1984532
Sequence 9285	AA333116	EST	1985360
Sequence 9286	AA335696	EST	1987937
Sequence 9287	AA335862	EST	1988104
Sequence 9288	AA336878	EST	1989117
Sequence 9289	AA337404	EST	1989869
Sequence 9290	AA337798	EST	1990057
Sequence 9291	AA338634	EST	1990882
Sequence 9292	AA339622	EST	1991880
Sequence 9293	AA341987	EST	1994222
Sequence 9294	AA343749	EST	1995986
Sequence 9295	AA349886	EST	2002205
Sequence 9296	AA349983	EST	2002303
Sequence 9297	AA351222	EST	2003561
Sequence 9298	AA352688	EST	2005008
Sequence 9299	AA353179	EST	2005499
Sequence 9300	AA353913	EST	2006231
Sequence 9301	AA355954	EST	2008272
Sequence 9302	AA355956	EST	2008274

Table 3-3

Sequence 9303	AA356182	EST	2008511
Sequence 9304	AA356195	EST	2008524
Sequence 9305	AA358978	EST	2011296
Sequence 9306	AA359611	EST	2012169
Sequence 9307	AA359735	EST	2012052
Sequence 9308	AA364033	EST	2016351
Sequence 9309	AA367068	EST	2019406
Sequence 9310	AA368087	EST	2020446
Sequence 9311	AA369721	EST	
Sequence 9312	AA370428	EST	2022039
Sequence 9313	AA371420		2022933
Sequence 9314	AA371957	EST	2023780
		EST	2024275
Sequence 9315 Sequence 9316	AA372674 AA375636	EST	2025182
Sequence 9317	AA376361	EST EST	2027956
			2028679
Sequence 9318	AA376458	EST	2028777
Sequence 9319	AA378378	EST	2030695
Sequence 9320	AA378849	EST	2031168
Sequence 9321	AA386348	EST	2038701
Sequence 9322	AA397884	EST	2050623 <sup>-</sup>
Sequence 9323	AA399142	EST	2052896
Sequence 9324	AA399177	EST	2052913
Sequence 9325	AA401252	EST	2055141
Sequence 9326	AA401261	EST	2055168
Sequence 9327	AA401526	EST	2053804
Sequence 9328	AA402063	EST	2056854
Sequence 9329	AA403047	EST	2055609
Sequence 9330	AA406425	EST	2064410
Sequence 9331	AA412426	EST	2071129
Sequence 9332	AA417300	EST	2077408
Sequence 9333	AA417999	EST	2079818
Sequence 9334	AA418045	EST	2079864
Sequence 9335	AA418096	EST	2079897
Sequence 9336	AA418249	EST	2080078
Sequence 9337	AA421643	EST	2100459
Sequence 9338	AA424317	EST	2103287
Sequence 9339	AA425387	EST	2106170
Sequence 9340	AA425619	EST	2106375
Sequence 9341	AA425658	EST	2106433
Sequence 9342	AA425751	EST	2106453
Sequence 9343	AA425761	EST	2106481
Sequence 9344	AA427920	EST	2112054
Sequence 9345	AA428948	EST	2110491
Sequence 9346 Sequence 9347	AA429871	EST	2113108
	AA431577	EST	2115285
Sequence 9348	AA433865	EST	2138779
Sequence 9349	AA436966	EST	2141880
Sequence 9350	AA442297	EST	2154175
Sequence 9351	AA442687	EST	2154565
Sequence 9352	AA446391	EST	2159056
Sequence 9353	AA448292	EST	2161962
Sequence 9354	AA449483	EST	2163233
Sequence 9355	AA449501	EST	2163251
Sequence 9356	AA449646	EST	2163396
Sequence 9357	AA450096	EST	2163846
Sequence 9358	AA450166	EST	2163916

Table 3-3

Sequence 93	59 AA	450276	EST	2164026
Sequence 93		452716	EST	2166385
Sequence 93		453565	EST	2167234
Sequence 93		453899	EST	2167568
Sequence 93		456454	EST	2179030
Sequence 93		457365	EST	2180085
Sequence 93		458613		
Sequence 93			EST	2183520
_ '		459167 464383	EST	2184074
Sequence 93		461383	EST	2186503
Sequence 93		463367 465460	EST	2188251
Sequence 93		465402	EST	2191569
Sequence 93		469033	EST	2195567
Sequence 93		470067	EST	2197376
Sequence 93		176876	EST	2205087
Sequence 93		178162	EST	2206796
Sequence 93		179499	EST	2208055
Sequence 93		180816	EST	2210368
Sequence 93		181233	EST	2210785
Sequence 93		181838	EST	2209516
Sequence 93	78 AA4	185255	EST	2214474
Sequence 93	79 AA4	185520	EST	2214739
Sequence 93	80 AA4	188447	EST	2215878
Sequence 93	81 AA4	188488	EST	2215919
Sequence 93	82 AA4	188513	EST	2215944
Sequence 93	83 AA4	189117	EST	2218719
Sequence 93	84 AA4	189288	EST	2218890
Sequence 93	85 AA4	190389	EST	2219562
Sequence 93	86 AA4	91213	EST	2220386
Sequence 93	87 AA4	191638	EST	2221200
Sequence 93		194334	EST	2224121
Sequence 93	89 AA4	194355	EST	. 2224142
Sequence 93		94387	EST	2224174
Sequence 93		95912	EST	2229233
Sequence 93		196486	EST	2229807
Sequence 93		01442	EST	2236409
Sequence 93		04269	EST	2240429
Sequence 93		05816	EST	224,1953
Sequence 93		07818	EST	2244257
Sequence 93		08038	EST	2244477
Sequence 93		14302	EST	2253810
Sequence 93		16531	EST	2256055
Sequence 94		22888	EST	2263600
Sequence 94		27374	EST	2269443
Sequence 94		32530	EST	2276784
Sequence 94		32673	EST	2276927
Sequence 94		33360	EST	2277456
Sequence 94		34734	EST	
Sequence 94		35663		2278987
•		35845	EST	2279916
Sequence 944 Sequence 944			EST	2280098
		52507	EST	2322761
Sequence 94		558357	EST	2328834
Sequence 94		65548	EST	2337187
Sequence 94		65867	EST	2337506
Sequence 94		68785	EST	2341839
Sequence 94		73427	EST	2347955
Sequence 94	14 AA5	74223	EST	<b>2348</b> 738

Table 3-3

Sequence 94	15 AA5	76016	EST	2350531
Sequence 94	16 AA5	76889	EST	2354363
Sequence 94		79611	EST	2357795
Sequence 94		81185	EST	2358957
Sequence 94		87264	EST	2398078
Sequence 94		<b>87</b> 570	EST	
•				2397245
Sequence 94		94562	EST	2409912
Sequence 94		98697	EST	2432369
Sequence 94		00084	EST	2433709
Sequence 94		00355	EST	2433980
Sequence 94		01289	EST	2434914
Sequence 94		<b>03</b> 876	EST	2437737
Sequence 94	27 AA6	04860	EST	2445724
Sequence 94		08613	EST	2457041
Sequence 94	29 AA6	09417	EST	2457845
Sequence 94	30 AA6	12668	EST	2463706
Sequence 94		25213	EST	2537598
Sequence 94		26629	EST	2539016
Sequence 94		27320	EST	2540337
Sequence 94		28491	EST	2540878
Sequence 94		28520	EST	2540907
Sequence 94		31993	EST	2555407
Sequence 94		34958	EST	2558172
Sequence 94		40716	EST	2565966
Sequence 94		41939	EST	
Sequence 94		<b>523</b> 88		2567157
Sequence 94			EST	2584040
•		59693	EST	2595847
Sequence 94		62024	EST	2616115
Sequence 94		63649	EST	2617640
Sequence 94		<b>688</b> 86	EST	2630385
Sequence 94		79400	EST	2659922
Sequence 94		79621	EST	2660143
Sequence 94		87221	EST	2675412
Sequence 94		87495	EST	2675686
Sequence 944		93932	EST	2694870
Sequence 94		94188	EST	2695126
Sequence 94		00208	EST	2703171
Sequence 94		01138	EST	2704303
Sequence 948		04533	EST	2714451
Sequence 94		05937	EST	2715855
Sequence 948		09002	EST	2718920
Sequence 945		16423	EST	2728697
Sequence 948	57 AA7.	22455	EST	2740162
Sequence 948	58 AA7:	22786	EST	2740493
Sequence 948	59 AA7	<b>30</b> 400	EST	2751604
Sequence 946	30 AA7	31394	EST	2753550
Sequence 946	31 AA7	42260	EST	2784260
Sequence 946	32 AA7	46310	EST	2786296
Sequence 946	33 AA7	<b>46</b> 748	EST	2786734
Sequence 946	34 AA7	47197	EST	2787155
Sequence 946	55 AA7	49387	EST	2789345
Sequence 946		57433	EST	2805296
Sequence 946		61101	EST	2810031
Sequence 946		65907	EST	2817145
Sequence 946		69588	EST	2820826
Sequence 947		72570	EST	2824353
4				202-000

Table 3-3

	7		
Sequence 9471	Ä <b>77</b> 4779	EST	2834113
Sequence 9472	AA775625	EST	2834959
Sequence 9473	AA805474	EST	2874224
Sequence 9474	AA808205	EST	2877611
Sequence 9475	AA808239	EST	2877645
Sequence 9476	AA813991	EST	2883587
Sequence 9477	AA814182	EST	2883778
Sequence 9478	AA814817	EST	
Sequence 9479	AA826906	EST	2884413
Sequence 9480	AA830072		2900903
Sequence 9481	AA831991	EST EST	2903171
Sequence 9482	AA833749		2905090
	AA837407	EST	2908517
Sequence 9483		EST	2912606
Sequence 9484	AA838157	EST	2913814
Sequence 9485	AA843207	EST	2929725
Sequence 9486	AA843238	EST	2929756
Sequence 9487	AA843878	EST	2930329
Sequence 9488	AA843908	EST	2930359
Sequence 9489	AA857964	EST	2946266
Sequence 9490	AA864964	EST	2959277
Sequence 9491	AA868452	EST	2963897
Sequence 9492	AA868722	EST	2964167
Sequence 9493	AA872507	EST	2968685
Sequence 9494	AA872913	EST	2969035
Sequence 9495	AA872991	EST	2969113
Sequence 9496	AA877350	EST	2986427
Sequence 9497	AA877789	EST	2986754
Sequence 9498	AA902103	EST	3037293
Sequence 9499	AA902974	EST	3038097
Sequence 9500	AA903253	EST	3038376
Sequence 9501	AA905330	EST	3040453
Sequence 9502	AA906868	EST	3042112
Sequence 9503	AA907186	EST	3042646
Sequence 9504	AA908460	EST	3047865
Sequence 9505	AA913203	EST	3052595
Sequence 9506	AA916003	EST	3055395
Sequence 9507	AA927612	EST	3077033
Sequence 9508	AA931566	EST	3085952
Sequence 9509	AA932278	EST	3086516
Sequence 9510	AA968524	EST	3143704
Sequence 9511	AA973790	EST	3148970
Sequence 9512	AA976114	EST	3151906
Sequence 9513	AA976653	EST	3154099
Sequence 9514	AA984122	EST	3162647
Sequence 9515	AA988106	EST	3173470
Sequence 9516	AA988525	EST	3174217
Sequence 9517	AA988576	EST	3174268
Sequence 9518	AA991580	EST	3178069
Sequence 9519	AF001542	EST	2529714
Sequence 9520	Al015791	EST	3230127
Sequence 9521	AI017095	EST	3231431
Sequence 9522	Al017902	EST	3232238
Sequence 9523	AI018625	EST	3233144
Sequence 9524	Al022209	EST	3239562
Sequence 9525	AI027035	EST	3244551
Sequence 9526	AI033991	EST	3254944

Table 3-3

Sequence 9527	AI034371	EST	3255324
Sequence 9528	AI040602	EST	3279796
Sequence 9529	AI051466	EST	3307000
Sequence 9530	AI052000	EST	3307991
Sequence 9531	AI056337	EST	3330203
Sequence 9532	AI056402	EST	3330268
Sequence 9533	AI061175	EST	
Sequence 9534	AI064910	EST	3336543
Sequence 9535	AI066583	EST	6359182
Sequence 9536	AI076630	EST	3367285
Sequence 9537	AI076953	EST	3405808
Sequence 9538	Al078217	EST	3404782
Sequence 9539	Al079510	EST	3412625
Sequence 9540	Al080497	EST	3415761
Sequence 9541	Al081780	EST	3416748
Sequence 9542	Al084794	EST	3418572
Sequence 9543	AI084916		3423217
Sequence 9544	Al085616	EST	3423339
Sequence 9545	AI088897	EST	3424039
Sequence 9546	Al090197	EST	3427956
Sequence 9547	Al090197 Al091173	EST	3429256
Sequence 9548	AI094514	EST	3430232
Sequence 9549	Al095450	EST	3433490
Sequence 9550	Al114692	EST	3434426
Sequence 9551	Al123418	EST	6360037
Sequence 9552	Al124792	EST EST	3539184
Sequence 9553	Al126005		3593306
Sequence 9554	Al129250	EST	3594519
Sequence 9555	AI133074	EST EST	3597764
Sequence 9556	AI133088	EST	6360390
Sequence 9557	Al133274	EST	6360404
Sequence 9558	Al133426	EST	6360590 6360742
Sequence 9559	Al139764	EST	3645736
Sequence 9560	Al142414	EST	3658773
Sequence 9561	AI146833	EST	3674515
Sequence 9562	Al148106	EST	3675788
Sequence 9563	AI150430	EST	3678899
Sequence 9564	Al151146	EST	3679615
Sequence 9565	AI159774	EST	3693133
Sequence 9566	AI161072	EST	3694377
Sequence 9567	AI168316	EST	3701486
Sequence 9568	Al168348	EST	3701518
Sequence 9569	AI186453	EST	3737091
Sequence 9570	AI187157	EST	3737795
Sequence 9571	AI189475	EST	3740684
Sequence 9572	Al193372	EST	3744581
Sequence 9573	AI198218	EST	3750824
Sequence 9574	Al199191	EST	3751797
Sequence 9575	Al199194	EST	3751800
Sequence 9576	Al202137	EST	3754743
Sequence 9577	Al203214	EST	3755820
Sequence 9578	Al207471	EST	6361479
Sequence 9579	Al207868	EST	3769810
Sequence 9580	AI214048	EST	3777649
Sequence 9581	Al214453	EST	3778054
Sequence 9582	AI216098	EST	3785139
		•	0.00103

Table 3-3

0500	11047046	EST	3797061
Sequence 9583	Al217246 Al244268	EST	3839665
Sequence 9584	Al248386	EST	3843783
Sequence 9585	Al249242	EST	3845771
Sequence 9586	Al251739	EST	3848268
Sequence 9587	Al261980	EST	3870183
Sequence 9588	Al261960 Al262251	EST	3870454
Sequence 9589		EST	3886329
Sequence 9590	AI267162	EST	3887015
Sequence 9591	Al267848 Al268335	EST	3887502
Sequence 9592	Al268362	EST	3887529
Sequence 9593	Al270685	EST	3889852
Sequence 9594	Al275466	EST	3897740
Sequence 9595	Al276933	EST	3899201
Sequence 9596	AI270933 AI277443	EST	3899711
Sequence 9597	Al277443 Al278328	EST	3900596
Sequence 9598	Al279625	EST	3917859
Sequence 9599	Al280561	EST	3918794
Sequence 9600	Al280301 Al281394	EST	3919627
Sequence 9601	Al283373	EST	3921606
Sequence 9602	Al283578	EST	3921811
Sequence 9603	Al283470	EST	3922715
Sequence 9604	Al290052	EST	3931718
Sequence 9605 Sequence 9606	Al290111	EST	3931777
Sequence 9607	Al290154	EST	3931820
Sequence 9608	Al291087	EST	3933861
Sequence 9609	Al292094	EST	3934868
Sequence 9610	Al298974	EST	3958628
Sequence 9611	Al333116	EST	4069675
Sequence 9612	Al334177	EST	4070736
Sequence 9613	Al334889	EST	4071816
Sequence 9614	Al337358	EST	4074285
Sequence 9615	Al339284	EST	4076211
Sequence 9616	Al341138	EST	4078065
Sequence 9617	Al341291	EST	4078218
Sequence 9618	Al342090	EST	4079017
Sequence 9619	Al343404	EST	4080610
Sequence 9620	Al344189	EST	4081395
Sequence 9621	Al348588	EST	4085794
Sequence 9622	Al349337	EST	4086543
Sequence 9623	Al349957	EST	4087163
Sequence 9624	Al350524	EST	4087730
Sequence 9625	Al352184	EST	4089390
Sequence 9626	Al355277	EST	4095430
Sequence 9627	Al356461	EST	4108082
Sequence 9628	Al357566	EST	4109187
Sequence 9629	Al357582	EST	4109203
Sequence 9630	A1360681	EST	4112302
Sequence 9631	Al361796	EST	4113417
Sequence 9632	Al373828	EST	4153694
Sequence 9633	Al378900	EST	4188753
Sequence 9634	A1379649	EST	4189502
Sequence 9635	Al381737	EST	4194518
Sequence 9636	Al381972	EST	4194753
Sequence 9637	Al393269	EST	4222816
Sequence 9638	Al420586	EST	4266517

Sequence 9639	Al435112	EST	4200000
Sequence 9640		EST	4300898
Sequence 9641		EST	4304204
Sequence 9642		EST	4305374
Sequence 9643		EST	4290157
Sequence 9644		EST	4307708
Sequence 9645			4310301
Sequence 9646		EST	4310833
Sequence 9647	Al458758	EST	4311178
Sequence 9648	Al460220	EST	4311337
Sequence 9649	AJ468419	EST	4313101
Sequence 9650	Al472084	EST	4330509
Sequence 9651		EST	4334174
Sequence 9652	Al472566	EST	4325611
Sequence 9653	A1479075	EST	4372243
Sequence 9654	Al480254	EST	4373422
Sequence 9655	AJ492032	EST	4393035
Sequence 9656	AJ494037	EST	4395040
Sequence 9657	AI525541	EST	4439676
Sequence 9658	AI525654	EST	4439789
Sequence 9659	Al538842	EST	4452977
Sequence 9660	AI557226	EST	4489589
	AI560973	EST	4511314
Sequence 9661 Sequence 9662	AI563921	EST	4522378
Sequence 9663	AI564719	EST	4523176
	AI570615	EST	4533989
Sequence 9664 Sequence 9665	AI580252	EST	4564628
Sequence 9666	Al580336 Al581098	EST	4564712
Sequence 9667	Al561098 Al597918	EST	4565474
Sequence 9668	Al608796	EST	4606977
Sequence 9669	A1609087	EST	4617963
Sequence 9670	Al610880	EST	4618254
Sequence 9671	Al623984	EST	4620047
Sequence 9672	Al624751	EST	4648915
Sequence 9673	Al628354	EST EST	4649682
Sequence 9674	Al628417	EST	4665154
Sequence 9675	Al632470	EST	4665217
Sequence 9676	Al633000	EST	4683800
Sequence 9677	Al637826	EST	4684330
Sequence 9678	Al638561	EST	4690060 4600705
Sequence 9679	AI640607	EST	4690795 4703716
Sequence 9680	AI650583	EST	4703716
Sequence 9681	Al651877	EST	4735856
Sequence 9682	Al672492	EST	4852223
Sequence 9683	Al684595	EST	4895889
Sequence 9684	AI685930	EST	4897224
Sequence 9685	Al690089	EST	4901383
Sequence 9686	AI692658	EST	4969998
Sequence 9687	Al694944	EST	4982844
Sequence 9688	Al697185	EST	4985085
Sequence 9689	AI720354	EST	5037610
Sequence 9690	Al720743	EST	5037610
Sequence 9691	A1738670	EST	5037999 5100651
Sequence 9692	AJ743182	EST	
Sequence 9693	Al743921	EST	5111470 5112209
Sequence 9694	AI750105	EST	
·		-01	5128369

Sequence 9695	Al751208	EST	5129472
Sequence 9696	AI751459	EST	5129723
Sequence 9697	AI751487	EST	5129751
Sequence 9698	AI752264	EST	5130528
Sequence 9699	AI752414	EST	5130678
Sequence 9700	A1752425	EST	5130689
Sequence 9701	A1760045	EST	5175712
Sequence 9702	AI761315	EST	5176982
Sequence 9703	AI765501	EST	5232010
Sequence 9704	AI766305	EST	5232814
Sequence 9705	AI767578	EST	5234087
Sequence 9706	Al767726	EST	5234235
Sequence 9707	Al792015	EST	5339731
Sequence 9708	Al795827	EST	5361287
Sequence 9709	AI797228	EST	5362700
Sequence 9710	AI800215	EST	5365687
Sequence 9711	AI806017	EST	5392583
Sequence 9712	A1807798	EST	5394364
Sequence 9713	AI808865	EST	5395431
Sequence 9714	AI810669	EST	5397235
Sequence 9715	AI814463	EST	5425678
Sequence 9716	AI814513	EST	5425728
Sequence 9717	AI815492	EST	5431038
Sequence 9718	AI816438	EST	5431984
Sequence 9719	AI818683	EST	5437762
Sequence 9720	AI819405	EST	543 <b>8484</b>
Sequence 9721	AI828075	EST	5448746
Sequence 9722	AI863382	EST	5527489
Sequence 9723	A1873001	EST	5547050
Sequence 9724	AI873501	EST	5547550
Sequence 9725	AI874326	EST	5548375
Sequence 9726	Al884491	EST	5589655
Sequence 9727	A1884783	EST	5589947
Sequence 9728	Al885021	EST	5590185
Sequence 9729	A1887758	EST	5592922
Sequence 9730	AI888327	EST	5593577
Sequence 9731	AI888872	EST	5594036
Sequence 9732	Al903705	EST	6494092
Sequence 9733	A1904447	EST	6494834
Sequence 9734	A1906328	EST	6496715
Sequence 9735	Al910090	EST	6500770
Sequence 9736	Al914133	EST	5633988
Sequence 9737	Al918605	EST	5638460
Sequence 9738	Al924534	EST	5660498
Sequence 9739	Al927755	EST	5663719
Sequence 9740	A1929696	EST	5665660
Sequence 9741	Al936166	EST	5675036
Sequence 9742	Al950991	EST	5743301
Sequence 9743	A1970095	EST	5766921
Sequence 9744	AI972423	EST	5769339
Sequence 9745	AI982834	EST	5810053
Sequence 9746	Al982890	EST	5810109
Sequence 9747	AL035972	EST	5405602
Sequence 9748	AL036220	EST	5405817
Sequence 9749	AL036415	EST	5405971
Sequence 9750	AL037386	EST	5406786

Table 3-3

Sequence 9751	AL037968	EST	5928298
Sequence 9752	AL039569	EST	5408608
Sequence 9753	AL039961	EST	5408942
Sequence 9754	AL040667	EST	5409613
Sequence 9755	AL040692	EST	5409638
Sequence 9756	AL041673	EST	5421022
Sequence 9757	AL042613	EST	5935505
Sequence 9758	AL042680	EST	5422129
Sequence 9759	AL042725	EST	5422174
Sequence 9760	AL042986	EST	5935655
Sequence 9761	AL043210	EST	5935803
Sequence 9762	AL043542	EST	5422929
Sequence 9763	AL043672	EST	5423059
Sequence 9764	AL045757	EST	5936079
Sequence 9765	AL045793	EST	5433905
Sequence 9766	AL045936	EST	5434039
Sequence 9767	AL045937	EST	5936123
Sequence 9768	AL045976	EST	5434076
Sequence 9769	AL046015	EST	5434109
Sequence 9770	AL047001	EST	5435057
Sequence 9771	AL047332	EST	4727279
Sequence 9772	AL047890	EST	4728078
Sequence 9773	AL048554	EST	5936565
Sequence 9774	AL110421	EST	5866029
Sequence 9775	AL118756	EST	5924655
Sequence 9776	AL118845	EST	5924744
Sequence 9777	AL119809	EST	5925708
Sequence 9778	AL120711	EST	5926610
Sequence 9779	AL120865	EST	5926866
Sequence 9780	AL120911	EST	5926912
Sequence 9781	AL121473	EST	5927474
Sequence 9782	AL121558	EST	5927559
Sequence 9783	AL133779	EST	6601967
Sequence 9784	AL133900	EST	6602087
Sequence 9785	AL134724	EST	6602911
Sequence 9786	AL134795	EST	6602982
Sequence 9787	AL134938	EST	6603125
Sequence 9788	AL135007	EST	6603194
Sequence 9789	AL135023	EST	6603210
Sequence 9790	AL135290	EST	6603477
Sequence 9791	AL135343	EST	6603530
Sequence 9792	AL135362	EST	6603549
Sequence 9793	AL135502	EST	6603689
Sequence 9794	AL135600	EST	6603787
Sequence 9795	AL138042	EST	6854722
Sequence 9796	AW002442	EST	5849358
Sequence 9797	AW014499	EST	5863256
Sequence 9798	AW014727	EST	5863484
Sequence 9799	AW029611	EST	5888367
Sequence 9800	AW029642	EST	5888398
Sequence 9801	AW043897	EST	5904426
Sequence 9802	AW062B37	EST	6014222
Sequence 9803	AW068104	EST	6023102
Sequence 9804	AW070431	EST	6025429
Sequence 9805 Sequence 9806	AW070690	EST	6025688
pedretine 2000	AW080672	EST	6035905

Table 3-3

Sequence 9807	AW084080	EST	6039232
Sequence 9808	AW118447	EST	6087031
Sequence 9809	AW148635	EST	6196531
•	AW149925	EST	6197821
Sequence 9810			
Sequence 9811	AW157231	EST	6228632
Sequence 9812	AW160468	EST	6299501
Sequence 9813	AW160833	EST	62 <b>99866</b>
Sequence 9814	AW161807	EST	63 <b>00840</b>
Sequence 9815	AW161978	EST	6301011
Sequence 9816	AW162587	EST	63 <b>01620</b>
Sequence 9817	AW162761	EST	6301794
Sequence 9818	AW162774	EST	6301807
Sequence 9819	AW163485	EST	6302518
Sequence 9820	AW163783	EST	6302816
Sequence 9821	AW175627	EST	6441768
	AW177931	EST	6443968
Sequence 9822			
Sequence 9823	AW178072	EST	6444109
Sequence 9824	AW194013	EST	6472742
Sequence 9825	AW237543	EST	6569932
Sequence 9826	AW239041	EST	6571431
Sequence 9827	AW239281	EST	6571 <b>6</b> 71
Sequence 9828	AW245631	EST	65 <b>88624</b>
Sequence 9829	AW245889	EST	65 <b>88882</b>
Sequence 9830	AW245906	EST	6588899
Sequence 9831	AW245923	EST	6588916
Sequence 9832	AW246049	EST	6589042
Sequence 9833	AW246074	EST	6589067
Sequence 9834	AW247459	EST	6590452
Sequence 9835	AW247494	EST	6590487
	AW249281	EST	6592274
Sequence 9836			6593118
Sequence 9837	AW250125	EST	
Sequence 9838	AW250126	EST	6593119
Sequence 9839	AW263239	EST	6640055
Sequence 9840	AW270779	EST	6657809
Sequence 9841	AW273055	EST	6660085
Sequence 9842	AW274583	EST	6661613
Sequence 9843	AW292896	EST	6699532
Sequence 9844	AW292897	EST	66 <b>99533</b>
Sequence 9845	AW293159	EST	6699795
Sequence 9846	AW294649	EST	67 <b>01285</b>
Sequence 9847	AW294971	EST	670 <b>1607</b>
Sequence 9848	AW296629	EST	67 <b>03265</b>
Sequence 9849	AW303456	EST	6713145
Sequence 9850	AW327374	EST	6797869
Sequence 9851	AW327483	EST	6797978
Sequence 9852	AW340172	EST	6836798
	AW340708	EST	6837334
Sequence 9853			
Sequence 9854	AW360965	EST	6865615
Sequence 9855	AW361001	E01	6865651
Sequence 9856	AW361103	EST	6865753
Sequence 9857	AW361711	EST	6866361
Sequence 9858	AW362982	EST	6867 <b>632</b>
Sequence 9859	AW363561	EST	68 <b>68211</b>
Sequence 9860	AW364748	EST	6869502
Sequence 9861	AW364827	EST	686 <b>9386</b>
Sequence 9862	AW365085	EST	68 <b>69735</b>
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Table 3-3

Sequence 9863	AW365852	EST	6870502
Sequence 9864	AW368293	EST	6872943
Sequence 9865	AW368461	EST	6873111
Sequence 9866	AW375589	EST	6880152
Sequence 9867	AW376917	EST	6881580
Sequence 9868	AW379827	EST	6884486
Sequence 9869	AW381227	EST	6885886
Sequence 9870	AW381248	EST	6885907
Sequence 9871	AW383599	EST	
Sequence 9872	AW384101	EST	688 <b>8167</b> 68 <b>8876</b> 0
Sequence 9873	AW384915	EST	
Sequence 9874	AW385192	EST	6889574
Sequence 9875	AW385322		6889851
Sequence 9876		EST	6889981
Sequence 9877	AW385328 AW388939	EST	6889987
•		EST	6893598
Sequence 9878	AW392421	EST	6897080
Sequence 9879	AW393095	EST	6897754
Sequence 9880	AW401915	EST	6920601
Sequence 9881	AW402136	EST	6920822
Sequence 9882	AW402676	EST	6921388
Sequence 9883	AW404132	EST	6923189
Sequence 9884	AW405567	EST	69 <b>24624</b>
Sequence 9885	AW407212	EST	6926269
Sequence 9886	AW408274	EST	6927331
Sequence 9887	AW408423	EST	6927480
Sequence 9888	AW410520	EST	69 <b>36061</b>
Sequence 9889	AW410772	EST	6936313
Sequence 9890	AW411211	EST	6936752
Sequence 9891	AW411217	EST	69 <b>36</b> 758
Sequence 9892	AW474433	EST	7044539
Sequence 9893	AW498895	EST	7137236
Sequence 9894	AW500117	EST	7112425
Sequence 9895	AW500758	EST	7113663
Sequence 9896	AW501342	EST	7114784
Sequence 9897	AW501373	EST	7114844
Sequence 9898	AW501390	EST	7114878
Sequence 9899	AW501401	EST	7114900
Sequence 9900	AW501421	EST	7114936
Sequence 9901	AW515416	EST	N/A
Sequence 9902	AW578462	EST	7253511
Sequence 9903	AW578582	EST	7253631
Sequence 9904	AW582604	EST	7257653
Sequence 9905	AW583629	EST	7260587
Sequence 9906	C04057	EST	1467308
Sequence 9907	F06849	EST	672465
Sequence 9908	F09531	EST	682064
Sequence 9909	F11462	EST	705760
Sequence 9910	F12434	EST	706790
Sequence 9911	H01873	EST	864806
Sequence 9912	H08043	EST	872865
Sequence 9913	H08093	EST	872915
Sequence 9914	H09228	EST	874050
Sequence 9915	H09317	EST	874139
Sequence 9916	H09793	EST	87 <b>4615</b>
Sequence 9917	H10561	EST	875383
Sequence 9918	H10893	EST	875713
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Table 3-3

Sequence 9919	H11277	EST	876097
Sequence 9920	H15211	EST	880031
Sequence 9921	H16865	EST	883105
Sequence 9922	H22726	EST	891421
Sequence 9923	H26047	EST	895170
Sequence 9924	H29998	EST	900908
Sequence 9925	H30612	EST	901522
Sequence 9926	H62143		
Sequence 9927	H69585	EST	1014975
Sequence 9928	H70297	EST	1039791
•		EST	1040503
Sequence 9929	H86644	EST	1068223
Sequence 9930	H99096	EST	1123764
Sequence 9931	M85706	EST	274353
Sequence 9932	N26971	EST	1141319
Sequence 9933	N28905	EST	1147141
Sequence 9934	N28944	EST	1147180
Sequence 9935	N34003	EST	1154403
Sequence 9936	N34221	EST	1155363
Sequence 9937	N36764	EST	1157906
Sequence 9938	N42722	EST	1167152
Sequence 9939	N43012	EST	1166756
Sequence 9940	N46795	EST	1187961
Sequence 9941	N54326	EST	1195646
Sequence 9942	N57253	EST	1201143
Sequence 9943	N95531	EST	1267800
Sequence 9944	R17855	EST	771465
Sequence 9945	R19920	EST	774554
Sequence 9946	R34894	EST	791795
Sequence 9947	R35355	EST .	792256
Sequence 9948	R35487	EST	792388
Sequence 9949	R50695	EST	812597
Sequence 9950	R52763	EST	814665
Sequence 9951	R53255	EST	815157
Sequence 9952	R59084	EST	829779
Sequence 9953	R69898	EST	843415
Sequence 9954	R73759	EST	848129
Sequence 9955	R84545	EST	942951
Sequence 9956	R88895	EST	953722
Sequence 9957	R89026	EST	953853
Sequence 9958	R93920	EST	969315
Sequence 9959	T05548	EST	316698
Sequence 9960	T08921	EST	389949
Sequence 9961	T11899	EST	596603
Seguence 9962	T18975	EST	601018
Sequence 9963	T19025	EST	601068
Sequence 9964	T32930	EST	615028
Sequence 9965	T63081	EST	666738
Sequence 9966	T77019	EST	694222
Sequence 9967	T84756	EST	713108
Sequence 9968	W02349	EST	
Sequence 9969	W03691	EST	1274356
Sequence 9970	W07619		1275555
Sequence 9971	W27890	EST	1281621
Sequence 9972		EST	1307838
Sequence 9973	W28867	EST	1308878
Sequence 9974	W38913	EST	1320619
Coquence 88/4	W38916	EST	1320622

Table 3-3

Sequence 9975	W39482	EST	1321190
Sequence 9976	W52153	· EST	1349763
Sequence 9977	W60983	EST	1367742
Sequence 9978	W69356	EST	1378667
Sequence 9979	W72044	EST	1382314
Sequence 9980	W76216	EST	1386441
Sequence 9981	W79237	EST	1390368
Sequence 9982	W80613	EST	1391700
Sequence 9983	W81613	EST	1392652
Sequence 9984	W88815	EST	1404287
Sequence 9985	W95032	EST	
Sequence 9986	Z25224	EST	1424152 395863
Sequence 9987			
	Z42427	EST	565844
Sequence 9988	Z45029	EST	574221
Sequence 9989	Z45988	EST	575222
Sequence 9990	Z78343	EST	1495116
Sequence 9991	Q46540	NUCPATENT	N/A
Sequence 9992	Q84560	NUCPATENT	N/A
Sequence 9993	T33624	NUCPATENT	N/A
Sequence 9994	T71086	NUCPATENT	N/A
Sequence 9995	T94119	NUCPATENT	N/A
Sequence 9996	V04699	NUCPATENT	N/A
Sequence 9997	V05728	NUCPATENT	N/A
Sequence 9998	V34175	NUCPATENT	N/A
Sequence 9999	V34185	NUCPATENT	N/A
Sequence 10000	V35457	NUCPATENT	N/A
Sequence 10001	V36303	NUCPATENT	N/A
Sequence 10002	V41451	NUCPATENT	N/A
Sequence 10003	V58631	NUCPATENT	N/A
Sequence 10004	V84411	NUCPATENT	N/A
Sequence 10005	V84546	NUCPATENT	N/A
Sequence 10006	V87058	NUCPATENT	N/A
Sequence 10007	V89685	NUCPATENT	N/A
Sequence 10008	V90259	NUCPATENT	N/A
Sequence 10009	V99828	NUCPATENT	N/A
Sequence 10010	X00616	NUCPATENT	N/A
Sequence 10011	X05715	NUCPATENT	N/A
Sequence 10012	X37479	NUCPATENT	N/A
Sequence 10013	X40381	NUCPATENT	N/A
Sequence 10014	X61430	NUCPATENT	N/A
Sequence 10015	X77342	NUCPATENT	N/A
Sequence 10016	X80672	NUCPATENT	N/A
Sequence 10017	X87868	NUCPATENT	N/A
Sequence 10018	X91484	NUCPATENT	N/A
Sequence 10019	X97576	NUCPATENT	N/A
Sequence 10020	X97718	NUCPATENT	N/A
Sequence 10021	X97772	NUCPATENT	N/A
Sequence 10022	X98309	NUCPATENT	N/A
Sequence 10023	X98400	NUCPATENT	N/A
Sequence 10024	X98781	NUCPATENT	N/A
Sequence 10025	X99117	NUCPATENT	N/A
Sequence 10026	Z00428	NUCPATENT	N/A
Sequence 10027	Z14314	NUCPATENT	N/A
Sequence 10028	Z14685	NUCPATENT	N/A
Sequence 10029	Z15023	NUCPATENT	N/A
Sequence 10030	Z24893	NUCPATENT	N/A
7-5.100 10000	EE-7000	HOULKIENI	13/74

Sequence 10031	Z27237	NUCPATENT	N/A
Sequence 10032	Z29638	NUCPATENT	N/A
Sequence 10033	Z38853	NUCPATENT	N/A
Sequence 10034	Z39903	NUCPATENT	N/A
Sequence 10035	Z41312	NUCPATENT	N/A
Sequence 10036	Z52414	NUCPATENT	N/A
Sequence 10037	Z52416	NUCPATENT	N/A
Sequence 10038	Z53013	NUCPATENT	N/A
Sequence 10039	Z57842	NUCPATENT	N/A
Sequence 10040	Z60383	NUCPATENT	N/A
Sequence 10041	Z65271	NUCPATENT	N/A
Sequence 10042	Z80193	NUCPATENT	N/A
Sequence 10043	Z80472	NUCPATENT	N/A
Sequence 10044	Z97065	NUCPATENT	N/A
Sequence 10045	Z97145	NUCPATENT	N/A
Sequence 10046	Z97322	NUCPATENT	N/A
Sequence 10047	AC05588	PREPATNUC	N/A
Sequence 10048	AC37772	PREPATNUC	N/A
Sequence 10049	AC38092	PREPATNUC	N/A
Sequence 10050	AC38120	PREPATNUC	N/A
Sequence 10051	AC38948	PREPATNUC	N/A
Sequence 10052	AC39251	PREPATNUC	N/A
Sequence 10053	AC39528	PREPATNUC	N/A
Sequence 10054	AC39561	PREPATNUC	N/A

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCTGTTGTGTGGCCTGGGCCGCAA
ACCCGATGTGGCCGAAAATTGGTAGGACCGTAGAACCCGATGTGCGA

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTCCACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGCCGCTGTTGTGTGGCCTGGGCCGCAA

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGÁTCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGCGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication W000/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGGTGGCAGTGTTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAAATAAAAAGGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication W000/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGCGCTGTTGTGTGGCCTGGGCCGCAA

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCGCTGTTGTGTGGCCTGGGCCGCAA

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

ANAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication W000/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAAATAAAAAGGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACCAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication W000/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGTGTTGTGTGGCCTGGGCCGCAA
A

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

ACTTGTGAGCCTGGACTGGCTTCCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT
CTCATCGAGGACTTGAGGCTGGCCTTGGAGATGCTGGAGCTTCCTCAGGAGAGAGCAGCC
CTCCTGAGCCAGATCCCTGGCCCAACAGCTGCCTACATAAAGGAATGGTTTGAAGAGAGC
CTCCTGAGCCAGGTAAGCCACCACAGTGCTGCTAGTAATGAAACCTACCAGGAACGCTTGGCA
CGTCTAGAAGGGGATAAGGAGTCCCTCATATTGCAGGTGAGTGTCCTCACAGACCAAGTA
GAAGCCCAGGGAGAAAAGATTCGAGACCTGGAAGTGTCTTGGAAGGACACCAGGTGAAA
CTCAATGCTGCTGAAGAGATGCTTCAACAGGAGCTGCTAAGCCGCACATCTCTTGAGACC
CAGAAGCTCGATCTGATGACTGAAGTGTCTGAGCTGAAGCTCAAGCTGGTTGGCATGAG
AAGGAGCAGAGAGAGAGAAAAGCAGAGAGAAAAGCAGAGAGTTACTGCAAGAGCTC
AGGCCCCTCAAAATCAAAGTGGAAGAGTTTGGAAA

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

### Table 3-4

ANAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication W000/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAATAAAAAGGAGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found in patent publication W000/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGCCC

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTGTCACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061
CCCGCGGTGGCCGAGGTACTGCGTGGCCGCAA

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT
CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT
TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA
Sequence 10053: Found in patent publication W000/06698
ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT
GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG
GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA
AGTTTGGGCCCCCTCCCTGCCCCAGCTTTCCTAGGCCAGAATTGTTTTGGCAGTAATTG
TCTGTTTAAAAAAATAAAAAGGAGGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA
TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG
CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC
GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCCCCATGCCTCACTGGTCCCA
GATCANGGCTTCAGGGACCAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT
ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTTGGAT
Sequence 10054: Found In patent publication W000/18922
TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGCCCCA

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTCACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGGCCGAGGTACTGCGTGGCCGTCGCTGTTGTGTGGCCTGGGCCGCAA

CCGGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

CCGCGGTGGCGGCCGAGGTCTGAACTGCGGGGTCTCTATCGCACTGCTAGGGGTTCTGCT GCTGGGTGCGGCGCGCGCGCGCGGGGCAGAAGCTTTTGAGATTGCTCTGCCACGAGA AAGCAACATTACAGTTCTCATAAAGCTGGGGACCCCGACTCTGCTGGCAAAACCCTGTTA CATCGTCTTTTCTAAAAGACATATAACCATGTTGTCCATCAAGTCTGGAGAAAAGAATAGT CTTTACCTTTAGCTGCCAGAGTCCTGAGAATCACTTTGTCATANAGATCCAGAAAAATAT TGACTGTATGTCAGGCCCATGTCCTTTTGGGGAGGTTCAGCTTCAGCCCTCGACATCGTT GTTGCCTACCCTCAACAGAACTTTCATCTGGGATGTCAAA

Sequence 10049: Found in patent publication W000/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730
ACTTGTGAGCCTGGACTGGCTTCCCGGCCTCCTACATGAACCCCTTCCCGGTGCTCCAT

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799
ACCGCGGGGCCGCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGT
CCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGT
CTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGAC
ATATCCAATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATT
TCATGTCCATTTTTAAATAATTCAAACATTTTGGGGATGTCACGGCCAATGGAATTTCGA
GAAAGCTTTGGATATTTTTTATTCAGTTTCTTCCTTAATCGATTAAGTTCAGGCATTATT
TCTGGAACAGCTACGTAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGA
CTAAGTGCCTCCTGCAAATGCAGCTCCATTTTCTTCCGCTATTTTGACCTCTGATGCATT
CTCTGTAAATACAGCAACTTTATTGATTTCCGAAGCCAATGGGTATGCAAACTAANAAC
ACTGGTAAATGGCTTCACGTTTTTCTTNTTTTCCAGTGCCATATNCAGTGTCAAAATCAA

#### Table 3-4

ANAAAACACTTTGCTTTGGACTAGTTAAAGCCAANAATTTGA Sequence 10053: Found in patent publication WO00/06698 ACATCTTCCAGGACAAGGTAAACTCTGACATGCACTAGGTATGTGCAGATCCCGGCCCCT GCCACCCAGCCTCATGCAAGTCATCCCCGACATGACCTTCACGACCGCAATGCAAGGAGG GGAAGAAAGTCACAGCACTGATGAGGACAGCTGCAGAGGTGGCAGTGTGTGGACACAGGA TCTGTTTAAAAAAATAAAAAGGAGGAGGAAGCGTTCACCGCCACAAATCATAAAATGGACA TGACTGTGGAGTCTTACAGTTCAGGGTTCTTTCATTCACGTCCCTTCCTGTCTCGGTCTG CGGTCTTTACCACATCAATAGGACTTTTTATGCGTCCGGGTTAATTTTTCACTCCAGTGC GTCCTGTTGCAGGGACCGGAGCTGATGGGAGCTGCTTCTCCCCATGCCTCACTGGTCCCA GATCANGGCTTCAGGGACAGATGATGAGTCTCAAACGAGCCANCAGGGGTTCTTTTGGGT ATAAATGGGCAATTCGNCCTGTCTTAAGNCTGATGACCTCANCCGTGGTTTTTGGAT Sequence 10054: Found in patent publication WO00/18922 TAGGGCGAATTGGAGCTCCCCGCGGTGGCGGCCGAGGTACCTGAGCCAGGAGGAGGCCC GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT GGCCGGGCTGAGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG GAGCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC TAACAAGCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC **ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC** 

#### <u>Table 3-4</u>

Sequence 10047: Found in patent publication WO99/09061 CCCGCGGTGCCGAGGTACTGCGTGCCGTCCGCTGTTGTGTGGCCTGGCCGCAA

CCGGCCACGTGGTTGACAATGGGGGAGAAATTGGTAGGACCGTAGAAGCGGATGTGGGG CAGGCAAGCTGAGTACGCCTGGGCAATACCATCCACACCTGAGCAGAAGGGGTTGGTGGG GTTGAAGTTGATGGCAAACTCATGGGAGACCTTCCAGTCTGGGGGTAACTGGGCCCCGAA TCCCAGAGCTGGAAACATCTTATCACTGTCGTAGTCCTGAATGATCTGCCCAACAGCCCA GATGGCCGACAGATATTCGTTGGTGCCCATAGGGTTGATATANTGCAAAGAGGAAGGGTC GAGGGGATTCCCGTTGGAGGCTGTAAAGTCTATTCCAACGGTGAACATGAGCTGGCAGCC TCCCAGGACGTAGTCAAGGAAGGAGTAGT

Sequence 10048: Found in patent publication WO00/06698

Sequence 10049: Found in patent publication WO00/06698

TGATCTTCAGCTGCTCCTGAACTTTGGCACTGTAAGGGTTCTCTGTATCGTATGGTGGTG ACAAAGCTTTTAAAATAACCTNTCCTCCAGTGCATGGAAAAGGACTGGAGTATTTATNAA GT

Sequence 10050: Found in patent publication WO00/06730

Sequence 10051: Found in patent publication WO00/11171

Sequence 10052: Found in patent publication WO00/15799

ACCGCGGGGGCCGNGGNTTCNGCCTGGAGGAGCTCAGGGTGGCCGGCATTCACAAG

AGGTGGCCCGGACCATCGGCATTTCTGTGGATCCGAGGAGGCGGAACAAGTCCACGGAGTCCCTGCAGACCAACGTGCAGCGGCTGAAGGAGTACCACAAAGGGACCCAAATTCAGCGGTCTGTGCCTACAAACTTCATTAATAACTGCTTGCAGATTGGCAGCTATCTGGTCACTTGACATATCCCATGTTGCTATTTTGGTCTGGAGAAAGTTCTCCCTTTCTTCATCTACCTTAATTTCATGCCATTTTTAAATAATTCAAACATTTTGGGGGATGTCACGGCCAATGGAATTTCGAGAACAGCTTTGGAACAGCTTCGTATCAGCATTATTTCTGGAACAGCTACGTAAAAAGTCTGCAACAATTTCATCATCCCAAATCTTCTGTATCAGACTAAGTGCCTCCTGAACAATTTCTTCCGCTATTTTTGACCTCTGATGCATTCCTCTGTAAAAACCTTTATTGATTTCCGAAGCCAATGGGTATGGCAAACTAAAAACAACTGGGTAAAATCAAAATCAAATCAAATCAAAATCAA

### Table 3-4

GGCCGTGGACCAGGAGCTATTTAACGAATACCAGTTCAGCGTGGACCAACTTATGGAACT
GGCCGGGCTGAGCTGTGCTACAGCCATCGCCAAGGCATATCCCCCCACGTCCATGTCCAG
GAGCCCCCCTACTGTCCTGGTCATCTGTGGCCCGGGGAATAATGGAGGAGATGGTCTGGT
CTGTGCTCGACACCTCAAACTCTTTGGCTACGAGCCAACCATCTATTACCCCAAAAGGCC
TAACAAGCCCCTCTTCACTGCATTGGTGACCCAGTGTCAGAAAATGGACATCCCTTTCCT
TGGGGAAATGCCCGCAGAGCCCATGACGATTGATGAACTGTTGAGCTGGTGGTGGATGCC
ATCTTTGGCTTCAGCTTCAAGGGCGATGTTCGGGAAC

# <u>Table 3-5</u>

Sequence #	Acc No	Database hit	GI Nbr
Sequence 10055	A02759	Genbank	345130
Sequence 10056	A36501	Genbank	2293812
Sequence 10057	A38056	Genbank	2294706
Sequence 10058	A46722	Genbank	2300822
	A98295		
Sequence 10059		Genbank	6781400
Sequence 10060	AB000584	Genbank	1813326
Sequence 10061	AB000888	Genbank	2467297
Sequence 10062	AB002323	Genbank	2224590
Sequence 10063	AB002337	Genbank	6683125
Sequence 10064	AB002383	Genbank	2224710
Sequence 10065	AB002533	Genbank	1944124
Sequence 10066	AB003102	Genbank	1945608
Sequence 10067	AB004064	Genbank	6983841
Sequence 10068	AB004885	Genbank	2217932
Sequence 10069	AB006534	Genbank	2924619
Sequence 10070	AB007170	Genbank	3077765
Sequence 10071	AB007956	Genbank	3413930
Sequence 10072	AB007964	Genbank	3413939
Sequence 10073	AB008226	Genbank	3370992
Sequence 10074	AB011004	Genbank	3273315
Sequence 10075	AB011083	Genbank	3043545
Sequence 10076	AB011100	Genbank	6683714
Sequence 10077	AB011120	Genbank	3043619
Sequence 10078	AB011165	Genbank	3043709
Sequence 10079	AB011399	Genbank	3452571
Sequence 10080	AB014511	Genbank	3327035
Sequence 10081	AB014570	Genbank	3327153
Sequence 10082	AB014579	Genbank	3327171
Sequence 10083	AB014583	Genbank	6635130
Sequence 10084	AB015639	Genbank	5821139
Sequence 10085	AB015798	Genbank	6681591
Sequence 10086	AB016089	Genbank	5821146
Sequence 10087	AB018257	Genbank	3882148
Sequence 10088	AB018274	Genbank	3882182
Sequence 10089	AB018284	Genbank	3882202
Sequence 10090	AB018304	Genbank	3882242
Sequence 10091	AB018330	Genbank	3882294
Sequence 10092	AB018340	Genbank	3882314
Sequence 10093	AB019564	Genbank	3885367
Sequence 10094	AB019568	Genbank	3885371
Sequence 10095	AB020335	Genbank	6518494
Sequence 10096	AB020676	Genbank	4240226
Sequence 10097	AB020683	Genbank	4240240
Sequence 10098	AB020694	Genbank	4240262
Sequence 10099	AB020864	Genbank	4003384
Sequence 10100	AB020867	Genbank	4003387
Sequence 10101	AB021179	Genbank	4062855
Sequence 10102	AB021467	Genbank	4586518
Sequence 10103	AB021663	Genbank	4996450
Sequence 10104	AB023050	Genbank	5672605
Sequence 10105	AB023051	Genbank	5672606
Sequence 10105	AB023140	Genbank	4589477
Sequence 10107	AB023140 AB023208	Genbank	
Sequence 10107 Sequence 10108	AB023206 AB024334		4589825
		Genbank	6016837
Sequence 10109	AB025355	Genbank	5103146

Sequence 10110	AB025959	Genbank	6520072
Sequence 10111	AB026894	Genbank	5821423
Sequence 10112	AB027466	Genbank	6172220
Sequence 10113	AB028624	Genbank	5103045
Sequence 10114	AB028859	Genbank	6567165
Sequence 10115	AB028893	Genbank	6552364
Sequence 10116	AB028975	Genbank	5689440
Sequence 10117	AB029001	Genbank	5689492
Sequence 10118	AB029025	Genbank	5689540
Sequence 10119	AB031052	Genbank	6978947
	AB031052 AB032957	Genbank	6329818
Sequence 10120			6329832
Sequence 10121	AB032959	Genbank	6330122
Sequence 10122	AB032982	Genbank	
Sequence 10123	AB033017	Genbank	6330344
Sequence 10124	AB033050	Genbank	6330623
Sequence 10125	AB033074	Genbank	6330846
Sequence 10126	AB033079	Genbank	6382025
Sequence 10127	AB034205	Genbank	6899845
Sequence 10128	AB034206	Genbank	6177737
Sequence 10129	AB035207	Genbank	6469033
Sequence 10130	AB037740	Genbank	7243018
Sequence 10131	AB037763	Genbank	7243064
Sequence 10132	AB037767	Genbank	7243072
Sequence 10133	AB037817	Genbank	7243172
Sequence 10134	AB037850	Genbank	7243238
Sequence 10135	AC000049	Genbank	5705995
Sequence 10136	AC000064	Genbank	1669369
Sequence 10137	AC000080	Genbank	7123907
Sequence 10138	AC000120	Genbank	1809224
Sequence 10139	AC000353	Genbank	6970735
Sequence 10140	AC000378	Genbank	2270906
Sequence 10141	AC000394	Genbank	2133911
Sequence 10142	AC001226	Genbank	2133862
Sequence 10143	AC001642	Genbank	1945597
Sequence 10144	AC002038	Genbank	2226439
Sequence 10145	AC002039	Genbank	2342716
Sequence 10145	AC002059 AC002051	Genbank	7798752
	AC002051	Genbank	2076723
Sequence 10147	AC002064 AC002347	Genbank	2828783
Sequence 10148		Genbank	
Sequence 10149	AC002350	Genbank	2961443
Sequence 10150	AC002357		2815516
Sequence 10151	AC002400	Genbank	2576344
Sequence 10152	AC002470	Genbank	6478944
Sequence 10153	AC002528	Genbank	2388554
Sequence 10154	AC002531	Genbank	2580476
Sequence 10155	AC002544	Genbank	3337382
Sequence 10156	AC002553	Genbank	3126783
Sequence 10157	AC002984	Genbank	2443872
Sequence 10158	AC003004	Genbank	2522284
Sequence 10159	AC003007	Genbank	2911728
Sequence 10160	AC003012	Genbank	2547258
Sequence 10161	AC003025	Genbank	3337308
Sequence 10162	AC003047	Genbank	3169149
Sequence 10163	AC003103	Genbank	2842782
Sequence 10164	AC003111	Genbank	2636670
Sequence 10165	AC003663	Genbank	3097871

## <u>Table 3-5</u>

Sequence 10166	AC003688	Genbank	3789719
Sequence 10167	AC003692	Genbank	2731601
Sequence 10168	AC003973	Genbank	2739352
Sequence 10169	AC004008	Genbank	2781388
Sequence 10170	AC004036	Genbank	2811104
Sequence 10171	AC004041	Genbank	2811099
Sequence 10172	AC004067	Genbank	3851208
Sequence 10173	AC004084	Genbank	2822156
Sequence 10174	AC004111	Genbank '	2828786
Sequence 10175	AC004148	Genbank	3482960
Sequence 10176	AC004151	Genbank	2896795
Sequence 10177	AC004131	Genbank	3386628
Sequence 10178	AC004218	Genbank	
Sequence 10179	AC004236	Genbank	4263838
Sequence 10180	AC004258	Genbank	2914668
Sequence 10181	AC004238	1	2924759
Sequence 10182	AC004388	Genbank	3046272
Sequence 10183	AC004368 AC004453	Genbank	3046271
Sequence 10184		Genbank	2979600
Sequence 10185	AC004496	Genbank	2996645
Sequence 10186	AC004527	Genbank	4760422
Sequence 10187	AC004530	Genbank	3108050
	AC004551	Genbank	3309005
Sequence 10188	AC004582	Genbank	3337307
Sequence 10189	AC004584	Genbank	3417305
Sequence 10190	AC004596	Genbank	4210521
Sequence 10191	AC004637	Genbank	3094998
Sequence 10192	AC004668	Genbank	3115345
Sequence 10193	AC004682	Genbank	3 <b>337388</b>
Sequence 10194	AC004687	Genbank	3 <b>258613</b>
Sequence 10195	AC004707	Genbank	32 <b>49127</b>
Sequence 10196	AC004742	Genbank	31 <b>52630</b>
Sequence 10197	AC004757	Genbank	3341707
Sequence 10198	AC004810	Genbank	3264567
Sequence 10199	AC004816	Genbank	4156209
Sequence 10200	AC004821	Genbank	4753291
Sequence 10201	AC004832	Genbank	6624129
Sequence 10202	AC004835	Genbank	4508154
Sequence 10203	AC004854	Genbank	4827328
Sequence 10204	AC004865	Genbank	4156195
Sequence 10205	AC004878	Genbank	3213119
Sequence 10206	AC004884	Genbank	4156189
Sequence 10207	AC004904	Genbank	4156180
Sequence 10208	AC004908	Genbank	4156179
Sequence 10209	AC004913	Genbank	3213084
Sequence 10210	AC004918	Genbank	4153871
Sequence 10211	AC004935	Genbank	4156170
Sequence 10212	AC004938	Genbank	3213059
Sequence 10213	AC004943	Genbank	3924671
Sequence 10214	AC004955	Genbank	4753267
Sequence 10215	AC004968	Genbank	4156160
Sequence 10216	AC004969	Genbank	4156159
Sequence 10217	AC004982	Genbank	3419846
Sequence 10218	AC004983	Genbank	4309885
Sequence 10219	AC004985	Genbank	5708490
Sequence 10220	AC004999	Genbank	3970963
Sequence 10221	AC005023	Genbank	3900847
			0000077

	•		
Sequence 10222	AC005035	Genbank	3953506
Sequence 10223	AC005041	Genbank	4508118
Sequence 10224	AC005057	Genbank	6587915
Sequence 10225	AC005058	Genbank	4156135
Sequence 10226	AC005069	Genbank	4753224
Sequence 10227	AC005070	Genbank	3406049
Sequence 10228	AC005071	Genbank	4508112
Sequence 10229	AC005095	Genbank	4753220
Sequence 10230	AC005101	Genbank	4508108
	AC005151	Genbank	3242763
Sequence 10231	AC005180	Genbank	3687281
Sequence 10232	AC005180 AC005189	Genbank	3264580
Sequence 10233		Genbank	3309101
Sequence 10234	AC005193	Genbank	3273381
Sequence 10235	AC005203	Genbank	3287444
Sequence 10236	AC005212	•	3282162
Sequence 10237	AC005218	Genbank	
Sequence 10238	AC005258	Genbank	3289992
Sequence 10239	AC005263	Genbank	3289978
Sequence 10240	AC005300	Genbank	6492490
Sequence 10241	AC005316	Genbank	3738342
Sequence 10242	AC005318	Genbank	3885345
Sequence 10243	AC005325	Genbank	3366581
Sequence 10244	AC005343	Genbank	4165008
Sequence 10245	AC005346	Genbank	3366566
Sequence 10246	AC005368	Genbank	3367506
Sequence 10247	AC005378	Genbank	4508139
Sequence 10248	AC005412	Genbank	5931398
Sequence 10249	AC005480	Genbank	4835818
Sequence 10250	AC005488	Genbank	5836194
Sequence 10251	AC005520	Genbank	5001541
Sequence 10252	AC005531	Genbank	4153875
Sequence 10253	AC005550	Genbank	3478668
Sequence 10254	AC005606	Genba <b>nk</b>	3540172
Sequence 10255	AC005612	Genbank	3540153
Sequence 10256	AC005618	Genbank	3548785
Sequence 10257	AC005630	Genbank	4159882
Sequence 10258	AC005661	Genbank	3818348
Sequence 10259	AC005740	Genbank	3687210
Sequence 10260	AC005754	Genbank	3688074
Sequence 10261	AC005786	Genbank	3702287
Sequence 10262	AC005789	Genbank	3702281
Sequence 10263	AC005799	Genbank	3789715
Sequence 10264	AC005828	Genbank	3789713
Sequence 10265	AC005876	Genbank	6249672
Sequence 10266	AC005971	Genbank	4314421
Sequence 10267	AC006001	Genbank	5708496
Sequence 10268	AC006008	Genbank	5091655
Sequence 10269	AC006017	Genbank	4508141
Sequence 10270	AC006040	Genbank	4314426
Sequence 10270	AC006055	Genbank	4071011
Sequence 10271	AC006057	Genbank	4731048
	AC006059	Genbank	4544348
Sequence 10273	AC006059 AC006064	Genbank	4572650
Sequence 10274	AC006064 AC006116	Genbank	3962497
Sequence 10275	AC006116 AC006160		5701616
Sequence 10276		Genbank	
Sequence 10277	AC006254	Genbank	4726099

Sequence 10278	AC006255	Genbank	5306223
Sequence 10279	AC006257	Genbank	4092478
Sequence 10280	AC006261	Genbank	4079613
Sequence 10281	AC006312	Genbank	4582483
Sequence 10282	AC006347	Genbank	4309919
Sequence 10283	AC006389	Genbank	4309883
Sequence 10284	AC006445	Genbank	6721147
Sequence 10285	AC006454	Genbank	5732182
Sequence 10286	AC006461	Genbank	4508128
Sequence 10287	AC006477	Genbank	7243871
Sequence 10287	AC006485	Genbank	4206211
	AC006485 AC006487	Genbank	4314418
Sequence 10289	AC006501	Genbank	4309874
Sequence 10290		Genbank	4220442
Sequence 10291	AC006504	Genbank Genbank	5815507
Sequence 10292	AC006509	- · ·	-
Sequence 10293	AC006518	Genbank	4713939
Sequence 10294	AC006543	Genbank	4388738
Sequence 10295	AC006561	Genbank	4558534
Sequence 10296	AC006581	Genbank	4914350
Sequence 10297	AC006960	Genbank	4337211
Sequence 10298	AC007041	Genbank	5708471
Sequence 10299	AC007055	Genbank	4885691
Sequence 10300	AC007191	Genbank	4558643
Sequence 10301	AC007204	Genbank	4559317
Sequence 10302	AC007225	Genbank	6715703
Sequence 10303	AC007226	Genbank	6715702
Sequence 10304	AC007227	Genbank	6456148
Sequence 10305	AC007325	Genbank	7229741
Sequence 10306	AC007528	Genbank	5263309
Sequence 10307	AC007546	Genbank	5668756
Sequence 10308	AC007637	Genbank	8439781
Sequence 10309	AC007684	Genbank	5836173
Sequence 10310	AC007708	Genbank	5923689
Sequence 10311	AC007790	Genbank	5042474
Sequence 10312	AC007842	Genbank	5080755
Sequence 10313	AC007967	Genbank	5931455
Sequence 10314	AC008082	Genbank	6143846
Sequence 10315	AC008101	Genbank	6970743
Sequence 10316	AC008116	Genbank	6001959
Sequence 10317	AC008124	Genbank	5815491
Sequence 10318	AC008179	Genbank	5931440
Sequence 10319	AC008269	Genbank	7243890
Sequence 10320	AC008498	Genbank	6850299
Sequence 10321	AC008893	Genbank	7259691
Sequence 10322	AC009363	Genbank	6648145
Sequence 10323	AC009509	Genbank	6492473
Sequence 10324	AC009946	Genbank	6604542
Sequence 10325	AC011331	Genbank	9929686
Sequence 10326	AC011362	Genbank	6094545
Sequence 10327	AC015853	Genbank	6721261
Sequence 10328	AC020663	Genbank	6682593
Sequence 10329	AD000092	Genbank	1905905
Sequence 10330	AE000658	Genbank	2358019
Sequence 10331	AF000160	Genbank	2306783
Sequence 10332	AF001437	Genbank	2316039
Sequence 10333	AF001549	Genbank	3355302
33400108 10003	A1 00 10-9	Goribank	0000002

<u>Table 3-5</u>

Sequence 10334	AF003001	Genbank	252944 <b>3</b>
Sequence 10335	AF005888	Genbank	<b>2</b> 738 <b>487</b>
Sequence 10336	AF006043	Genbank	2674061
Sequence 10337	AF006088	Genbank	2282041
Sequence 10338	AF006501	Genbank	4836840
			2852 <b>616</b>
Sequence 10339	AF007140	Genbank	
Sequence 10340	AF007544	Genbank	2970122
Sequence 10341	AF010238	Genbank	2282063
Sequence 10342	AF011390	Genbank	3298567
Sequence 10343	AF012072	Genbank	99675 <b>56</b>
Sequence 10344	AF014807	Genbank	2338731
Sequence 10345	AF014955	Genbank	2772828
Sequence 10346	AF015812	Genbank	25993 <b>59</b>
Sequence 10347	AF020351	Genbank	2655052
Sequence 10348	AF020352	Genbank	2655054
Sequence 10349	AF021819	Genbank	2460317
		Genbank	2465723
Sequence 10350	AF022211		
Sequence 10351	AF026126	Genbank	2828601
Sequence 10352	AF026292	Genbank	2559009
Sequence 10353	AF028832	Genbank	32874 <b>88</b>
Sequence 10354	AF029786	Genbank	3403166
Sequence 10355	AF031416	Genbank	3213216
Sequence 10356	AF032387	Genbank	26415 <b>56</b>
Sequence 10357	AF033095	Genbank	2645 <b>728</b>
Sequence 10358	AF034759	Genbank	2653648
Sequence 10359	AF035286	Genbank	2661038
Sequence 10360	AF035295	Genbank	2661050
Sequence 10361	AF035313	Genbank	2661075
	AF037222	Genbank	2921499
Sequence 10362			
Sequence 10363	AF037447	Genbank	6466790
Sequence 10364	AF037448	Genbank	3037012
Sequence 10365	AF038196	Genbank	2795917
Sequence 10366	AF038200	Genbank	2795 <b>921</b>
Sequence 10367	AF038554	Genbank	<b>2</b> 8955 <b>58</b>
Sequence 10368	AF038954	Genbank	<b>3</b> 32 <b>9377</b>
Sequence 10369	AF038955	Genbank	3329379
Sequence 10370	AF038958	Genbank	3329385
Sequence 10371	AF038966	Genbank	2791681
Sequence 10372	AF039029	Genbank	3834389
Sequence 10373	AF042836	Genbank	3776241
Sequence 10374	AF042857	Genbank	2952271
Sequence 10375	AF044956	Genbank	5326827
Sequence 10376	AF044957	Genbank	4164445
Sequence 10376	AF044959	Genbank	33481 <b>36</b>
Sequence 10377			
Sequence 10378	AF047181	Genbank	2909853
Sequence 10379	AF047431	Genbank	5514630
Sequence 10380	AF047436	Genbank	3335127
Sequence 10381	AF047439	Genbank	33351 <b>33</b>
Sequence 10382	AF047440	Genbank	<b>3</b> 3351 <b>35</b>
Sequence 10383	AF048686	Genbank	<b>5737698</b>
Sequence 10384	AF049140	Genbank	2947300
Sequence 10385	AF049613	Genbank	3329428
Sequence 10386	AF050637	Genbank	4164451
Sequence 10387	AF050638	Genbank	5326819
Sequence 10388	AF050639	Genbank	4164453
Sequence 10389	AF051894	Genbank	3095110

**Table 3-5** 

_			
Sequence 10390	AF052101	Genbank	3360408
Sequence 10391	AF052129	Genbank	3360438
Sequence 10392	AF052642	Genbank	2981453
Sequence 10393	AF053356	Genbank	3135305
Sequence 10394	AF053551	Genbank	3283048
Sequence 10395	AF053719	Genbank	6670884
Sequence 10396	AF054187	Genbank	4092059
Sequence 10397	AF054990	Genbank	3005703
Sequence 10398	AF054996	Genbank	3005711
Sequence 10399	AF055029	Genbank	3005759
Sequence 10400	AF055460	Genbank	3335143
Sequence 10401	AF055473	Genbank	nil
Sequence 10402	AF063020	Genbank	3283351
Sequence 10403	AF064019	Genbank	3347856
Sequence 10404	AF064607	Genbank	3152667
Sequence 10405	AF067518	Genbank	3850315
Sequence 10406	AF067656	Genbank	3901271
Sequence 10407	AF067844	Genbank	4240386
Sequence 10408	AF068846	Genbank	3201999
Sequence 10409	AF069378	Genbank	3982733
Sequence 10410	AF070598	Genbank	
Sequence 10411	AF070635	Genbank	3387976 3283905
Sequence 10412	AF070644	Genbank	3283917
Sequence 10413	AF070649	Genbank	
Sequence 10414	AF070655	Genbank	3283923
Sequence 10415	AF070659	Genbank	4454685
Sequence 10416	AF073298	Genbank	4454693
Sequence 10417	AF077030	Genbank	3641537
Sequence 10418	AF077036	Genbank	4689107 4689119
Sequence 10419	AF077037	Genbank	4689121
Sequence 10420	AF077043	Genbank	4689133
Sequence 10421	AF077202	Genbank	4679017
Sequence 10422	AF080092	Genbank	4322311
Sequence 10423	AF081484	Genbank	3420928
Sequence 10424	AF082513	Genbank	3649656
Sequence 10425	AF083441	Genbank	5813822
Sequence 10426	AF085357	Genbank	5114048
Sequence 10427	AF085359	Genbank	5114052
Sequence 10428	AF085867	Genbank	3483188
Sequence 10429	AF086090	Genbank	3483435
Sequence 10430	AF086207	Genbank	3483552
Sequence 10431	AF086210	Genbank	3483555
Sequence 10432	AF086234	Genbank	3483579
Sequence 10433	AF086378	Genbank	3483723
Sequence 10434	AF086388	Genbank	3483733
Sequence 10435	AF086557	Genbank	3483902
Sequence 10436	AF087481	Genbank	4322487
Sequence 10437	AF087659	Genbank	4191345
Sequence 10438	AF088028	Genbank	3523234
Sequence 10439	AF088036	Genbank	3523234
Sequence 10440	AF091071	Genbank	3859979
Sequence 10441	AF092094	Genbank	4426606
Sequence 10442	AF092130	Genbank	5138909
Sequence 10443	AF092135	Genbank	5138919
Sequence 10444	AF094481	Genbank	
Sequence 10445	AF097026	Genbank	4140681
	001020	Genbank	6409120

<u>Table 3-5</u>

0	AF097514	Genbank	4808600
Sequence 10446	AF097514 AF098162	Genbank	3929582
Sequence 10447	AF100928	Genbank	4323586
Sequence 10448	AF100928 AF105278	Genbank	4324471
Sequence 10449	AF106622	Genbank	4378528
Sequence 10450	AF108022 AF109134	Genbank	4139227
Sequence 10451	AF109154 AF109161	Genbank	4193945
Sequence 10452	AF110460	Genbank	4324698
Sequence 10453	AF111713	Genbank	5326796
Sequence 10454	AF111715	Genbank	6563217
Sequence 10455	AF112218	Genbank	6563223
Sequence 10456	AF112222	Genbank	6563229
Sequence 10457	AF113016	Genbank	6642755
Sequence 10458	AF113700	Genbank	6855634
Sequence 10459	AF117231	Genbank	6563235
Sequence 10460	AF117616	Genbank	4886909
Sequence 10461	AF118091	Genbank	6650827
Sequence 10462	AF121219	Genbank	4903275
Sequence 10463	AF121863	Genbank	4689265
Sequence 10464	AF121003	Genbank	5059061
Sequence 10465	AF125100	Genbank	5106995
Sequence 10466	AF125100 AF125102	Genbank	5106999
Sequence 10467	AF126021	Genbank	6563273
Sequence 10468	AF126782	Genbank	6318547
Sequence 10469	AF126762 AF126962	Genbank	6469610
Sequence 10470	AF128527	Genbank	4928043
Sequence 10471	AF120327	Genbank	4337095
Sequence 10472	AF129730	Genbank	4558506
Sequence 10473	AF131738	Genbank	4406548
Sequence 10474	AF131838	Genbank	4406677
Sequence 10475	AF131857	Genbank	4406704
Sequence 10476 Sequence 10477	AF132938	Genbank	4680646
Sequence 10477	AF132939	Genbank	4680648
Sequence 10479	AF132956	Genbank	4680682
Sequence 10479	AF132973	Genbank	4680716
Sequence 10481	AF132984	Genbank	4959567
Sequence 10482	AF134726	Genbank	4529886
Sequence 10483	AF144755	Genbank	5006628
Sequence 10484	AF146651	Genbank	5020073
Sequence 10485	AF147330	Genbank	4761681
Sequence 10486	AF151028	Genbank	7106777
Sequence 10487	AF151047	Genbank	7106815
Sequence 10488	AF151109	Genbank	6166337
Sequence 10489	AF151807	Genbank	4929566
Sequence 10490	AF151832	Genbank	4929616
Sequence 10491	AF151846	Genbank	4929644
Sequence 10492	AF151861	Genbank	4929674
Sequence 10493	AF151869	Genbank	4929690
Sequence 10494	AF151887	Genbank	4929726
Sequence 10495	AF151905	Genbank	4929762
Sequence 10495	AF152330	Genbank	5456956
Sequence 10497	AF153612	Genbank	4929830
Sequence 10498	AF156965	Genbank	5731112
Sequence 10499	AF161410	Genbank	6841233
Sequence 10499	AF161421	Genbank	6841255
Sequence 10501	AF161425	Genbank	6841263
Ochretice (000)	,		

<u>Table 3-5</u>

Sequence 10502	AF161428	Genbank	6841269
Sequence 10503	AF161448	Genbank	6841309
Sequence 10504	AF161485	Genbank	6841493
Sequence 10505	AF161490	Genbank	6841503
Sequence 10506	AF161503	Genbank	6841529
Sequence 10507	AF161556	Genbank	6841379
Sequence 10508	AF161800	Genbank	
Sequence 10509	AF167570	Genbank	5353769
Sequence 10510	AF168956	Genbank	5762314
Sequence 10511	AF172368	Genbank	5702387
Sequence 10512	AF176796	Genbank	5679370 5759308
Sequence 10513	AF179274	Genbank	5911402
Sequence 10514	AF180473	Genbank	6856202
Sequence 10515	AF186249	Genbank	6572947
Sequence 10516	AF187554	Genbank	6653225
Sequence 10517	AF190465	Genbank	6120105
Sequence 10518	AF196779	Genbank	6180170
Sequence 10519	AF199339	Genbank	6708280
Sequence 10520	AF203815	Genbank	6979641
Sequence 10521	AF207664	Genbank	6685071
Sequence 10522	AF213884	Genbank	7012904
Sequence 10523	AF217403	Genbank	6691160
Sequence 10524	AJ000519	Genbank	2739214
Sequence 10525	AJ000881	Genbank	2924308
Sequence 10526	AJ001443	Genbank	6006514
Sequence 10527	AJ001866	Genbank	2969902
Sequence 10528	AJ001981	Genbank	2546978
Sequence 10529	AJ002030	Genbank	2570006
Sequence 10530	AJ006291	Genbank	3805937
Sequence 10531	AJ010069	Genbank	3483012
Sequence 10532 Sequence 10533	AJ010597	Genbank	3559873
Sequence 10533	AJ011497	Genbank	4128014
Sequence 10535	AJ011930	Genbank	3859769
Sequence 10536	AJ012409	Genbank	3881975
Sequence 10537	AJ133812 AJ223352	Genbank	4704217
Sequence 10538	AJ224442	Genbank	3255996
Sequence 10539	AJ245719	Genbank	2911586
Sequence 10540	AK000023	Genbank Genbank	6688168
Sequence 10541	AK000028	Genbank	7019837
Sequence 10542	AK000086	Genbank	7019844
Sequence 10543	AK000107	Genbank	7019944
Sequence 10544	AK000221	Genbank	7019979
Sequence 10545	AK000291	Genbank	7020163 7020274
Sequence 10546	AK000318	Genbank	7020274
Sequence 10547	AK000348	Genbank	7020322
Sequence 10548	AK000395	Genbank	7020373
Sequence 10549	AK000414	Genbank	7020486
Sequence 10550	AK000426	Genbank	7020505
Sequence 10551	AK000434	Genbank	7020520
Sequence 10552	AK000439	Genbank	7020528
Sequence 10553	AK000474	Genbank	7020586
Sequence 10554	AK000486	Genbank	7020607
Sequence 10555	AK000507	Genbank	7020644
Sequence 10556	AK000541	Genbank	7020705
Sequence 10557	AK000548	Genbank	7020717

Sequence 10558	AK000560	Genbank	7020738
Sequence 10559	AK000571	Genbank	7020756
Sequence 10560	AK000585	Genbank	7020779
Sequence 10561	AK000667	Genbank	7020906
Sequence 10562	AK000756	Genbank	7021041
Sequence 10563	AK000790	Genbank	7021092
Sequence 10564	AK000915	Genbank	7021878
Sequence 10565	AK000934	Genbank	7021909
Sequence 10566	AK001061	Genbank	7022095
Sequence 10567	AK001091	Genbank	7022140
Sequence 10568	AK001207	Genbank	7022315
Sequence 10569	AK001313	Genbank	7022490
Sequence 10570	AK001314	Genbank	7022491
Sequence 10571	AK001321	Genbank	7022503
Sequence 10572	AK001375	Genbank	7022596
Sequence 10573	AK001387	Genbank	7022615
Sequence 10574	AK001395	Genbank	7022625
Sequence 10575	AK001413	Genbank	7022655
Sequence 10576	AK001430	Genbank	7022682.
Sequence 10577	AK001441	Genbank	7022699
Sequence 10578	AK001448	Genbank	7022713
Sequence 10579	AK001451	Genbank	7022717
Sequence 10580	AK001452	Genbank	7022718
Sequence 10581	AK001529	Genbank	7022840
Sequence 10582	AK001571	Genbank	7022906
Sequence 10583	AK001576	Genbank	7022913
Sequence 10584	AK001607	Genbank	7022965
Sequence 10585	AK001614	Genbank	7022977
Sequence 10586	AK001691	Genbank	7023106
Sequence 10587	AK001732	Genbank	7023176
Sequence 10588	AK001755	Genbank	7023219
Sequence 10589	AK001762	Genbank	7023231
Sequence 10590	AK002036	Genbank	7023674
Sequence 10591	AK002072	Genbank	7023732
Sequence 10592	AL008725	Genbank	2791551
Sequence 10593	AL009181	Genbank	2853179
Sequence 10594	AL021368	Genbank	3080468
Sequence 10595	AL021391	Genbank	4467344
Sequence 10596	AL021807	Genbank	9367202
Sequence 10597	AL021938	Genbank	3242170
Sequence 10598	AL022069	Genbank	3256174
Sequence 10599	AL022240	Genbank	4826471
Sequence 10600	AL022312	Genbank	4914501
Sequence 10601	AL022313	Genbank	4200326
Sequence 10602	AL022328	Genbank	5263010
Sequence 10603	AL022394	Genbank	9581758
Sequence 10604	AL023656	Genbank	5679746
Sequence 10605	AL024474	Genbank	3395511
Sequence 10606	AL024493	Genbank	3288439
Sequence 10607	AL030996	Genbank	3688349
Sequence 10608	AL031178	Genbank	3550040
Sequence 10609	AL031390	Genbank	5002609
Sequence 10610	AL031652	Genbank	9368398
Sequence 10611	AL031659	Genbank	9581755
Sequence 10612	AL031670	Genbank	4469083
Sequence 10613	AL031685	Genbank	9368423

<u>Table 3-5</u>

Sequence 10614	AL033397	Genbank	4902626
Sequence 10615	AL033518	Genbank	5763714
Sequence 10616	AL033531	Genbank	6807582
Sequence 10617	AL034371	Genbank	4938279
Sequence 10618	AL034374	Genbank	4455565
Sequence 10619	AL034379	Genbank	5918013
Sequence 10620	AL034400	Genbank	4455461
Sequence 10621	AL034548	Genbank	7263904
Sequence 10622	AL034582	Genbank	5830348
Sequence 10623	AL035209	Genbank	4160217
Sequence 10624	AL035304	Genbank	4200231
Sequence 10625	AL035413	Genbank	
Sequence 10626	AL035413 AL035422	Genbank	6010110
Sequence 10627	AL035458	Genbank	5263001 6624641
Sequence 10628	AL035436 AL035541	Genbank	
Sequence 10629	AL035661	Genbank	9581764
Sequence 10630	AL035699		6015535
Sequence 10631	AL033699 AL049280	Genbank	4826515
Sequence 10632	AL049280 AL049442	Genbank	4500037
•	AL049442 AL049563	Genbank	4500222
Sequence 10633 Sequence 10634	AL049563 AL049610	Genbank	4902753
Sequence 10635	and the second s	Genbank	5679448
· · · · · · · · · · · · · · · · · · ·	AL049712	Genbank	5629919
Sequence 10636	AL049779	Genbank	8176894
Sequence 10637 Sequence 10638	AL049823	Genbank	6273553
_	AL049829	Genbank	8217859
Sequence 10639 Sequence 10640	AL049830	Genbank	6729561
	AL049835	Genbank	5708093
Sequence 10641 Sequence 10642	AL049843	Genbank	5830432
•	AL050161	Genbank	4884375
Sequence 10643	AL050265	Genbank	4886440
Sequence 10644	AL050275	Genbank	4886500
Sequence 10645	AL050331	Genbank	5668655
Sequence 10646	AL050343	Genbank	6137008
Sequence 10647 Sequence 10648	AL050363	Genbank	4914597
Sequence 10649	AL050383	Genbank	4914585
Sequence 10650	AL050392 AL078621	Genbank	4914613
Sequence 10650		Genbank	6013067
_ •	AL079295	Genbank	5102607
Sequence 10652 Sequence 10653	AL079312 AL079342	Genbank Genbank	5102890
Sequence 10654	AL080080	Genbank Genbank	6018784
Sequence 10655	AL080084		5262491
Sequence 10656	AL080085	Genbank Genbank	5262498
Sequence 10657	AL080088	Genbank	5262499
Sequence 10658	AL080089		5262504
Sequence 10659	AL080139	Genbank Genbank	5262506
Sequence 10660	AL080178	Genbank	5262583
Sequence 10661	AL080243		5262652
Sequence 10662	AL080276	Genbank	5870622
Sequence 10663	_	Genbank	5763753
Sequence 10664	AL080317 AL096701	Genbank	5830430
Sequence 10665	AL096764	Genbank	5912599
Sequence 10665		Genbank	7228319
Sequence 10667	AL109613	Genbank	6136931
Sequence 10668	AL109754	Genbank	6911635
Sequence 10669	AL109758	Genbank	8217865
gednetics 10009	AL109925	Genbank	7105935

Sequence 10670	AL110183	Genbank	5817095
Sequence 10671	AL110185	Genbank	5817098
Sequence 10672	AL110202	Genbank	5817121
Sequence 10673	AL117350	Genbank	7159817
Sequence 10674	AL117461	Genbank	5911922
Sequence 10675	AL117481	Genbank	5911958
Sequence 10676	AL117694	Genbank	8248721
Sequence 10677	AL121601	Genbank	7159760
Sequence 10678	AL121694	Genbank	6318179
Sequence 10679	AL121716	Genbank	7159797
	AL122003	Genbank	6580480
Sequence 10680		Genbank	7159786
Sequence 10681	AL132821		
Sequence 10682	AL132992	Genbank	6996086
Sequence 10683	AL133060	Genbank Genbank	6453483
Sequence 10684	AL133163		6562367
Sequence 10685	AL133216	Genbank	6983473
Sequence 10686	AL133245	Genbank	7159621
Sequence 10687	AL133396	Genbank	6562003
Sequence 10688	AL133585	Genbank	6599171
Sequence 10689	AL135744	Genbank	6682291
Sequence 10690	AL135784	Genbank	6911343
Sequence 10691	AL135858	Genbank	7159622
Sequence 10692	AL136059	Genbank	7106618
Sequence 10693	AL136296	Genbank	8217913
Sequence 10694	AL136543	Genbank	6807646
Sequence 10695	AL136593	Genbank	7018431
Sequence 10696	AL136884	Genbank	6807672
Sequence 10697	AL137229	Genbank	7009597
Sequence 10698	AL137440	Genbank	6808001
Sequence 10699	AL137489	Genbank	6808110
Sequence 10700	AL137496	Genbank	6808137
Sequence 10701	AL137519	Genbank	6808181
Sequence 10702	AL137543	Genbank	6808222
Sequence 10703	AL137686	Genbank	6807956
Sequence 10704	AL137692	Genbank	6808008
Sequence 10705	AL137751	Genbank	6808387
Sequence 10706	AL137818	Genbank	7009598
Sequence 10707	AP000046	Genbank	3132356
Sequence 10708	AP000085	Genbank	4730826
Sequence 10709	AP000217	Genbank	4827264
Sequence 10710	AP000345	Genbank	5103008
Sequence 10711	AP000350	Genbank	5103013
Sequence 10712	AP000352	Genbank	6016843
Sequence 10713		Genbank	5926687
Sequence 10714	AP000525	Genbank	5931503
Sequence 10715	AP000700	Genbank	6705908
Sequence 10716	AP000962	Genbank	6942330
Sequence 10717		Genbank	6693602
Sequence 10718		Genbank	6970360
Sequence 10719		Genbank	6970361
Sequence 10719		Genbank	7262569
Sequence 10721	AP001172	Genbank	6983884
Sequence 10721		Genbank	7077183
Sequence 10723		Genbank	7209831
		Genbank	7209834
Sequence 10724			
Sequence 10725	D13866	Genbank	433410

## <u>Table 3-5</u>

Sequence 10726	D13900	Genbank	433412
Sequence 10727	D14043	Genbank	219924
Sequence 10728	D14658	Genbank	285940
Sequence 10729	D14662	Genbank	285948
Sequence 10730	D14665	Genbank	6630617
Sequence 10731	D14696	Genbank	285962
Sequence 10732	D14710	Genbank	559324
Sequence 10733	D15057	Genbank	493244
Sequence 10734	D16111	Genbank	435637
Sequence 10735	D16234	Genbank	303617
Sequence 10736	D16481	Genbank	473711
Sequence 10737	D16938	Genbank	598868
Sequence 10738	D16947	Genbank	598932
Sequence 10739	D21089	Genbank	475156
Sequence 10740	D25218	Genbank	434778
Sequence 10741	D26600	Genbank	565650
Sequence 10742	D29805	Genbank	474986
Sequence 10743	D38112	Genbank	644480
Sequence 10744	D42041	Genbank	577294
Sequence 10745	D42052	Genbank	1136740
Sequence 10746	D42063	Genbank	924266
Seguence 10747	D45915	Genbank	1483130
Sequence 10748	D49355	Genbank .	1020405
Sequence 10749	D50372	Genbank	2605593
Sequence 10750	D50929	Genbank	1469200
Sequence 10751	D63486	Genbank	1469885
Sequence 10752	D63874	Genbank	968887
Sequence 10753	D66904	Genbank	1235726
Sequence 10754	D67025	Genbank	2656091
Sequence 10755	D79986	Genbank	1136389
Sequence 10756	D80000	Genbank	1136415
Sequence 10757	D80007	Genbank	1136429
Sequence 10758	D83253	Genbank	2951772
Sequence 10759	D84907	Genbank	1596053
Sequence 10760	D85181	Genbank	1906795
Sequence 10761	D86962	Genbank	1503997
Sequence 10762 Sequence 10763	D87018 D87127	Genbank	2114280
Sequence 10764	D87666	Genbank	1817551
Sequence 10765	D87742	Genbank Genbank	1620016
Sequence 10766	D89052	Genbank	1665824 1694672
Sequence 10767	E01716	Genbank	2169969
Sequence 10768	E01932	Genbank	2170180
Sequence 10769	E01954	Genbank	2170180
Sequence 10770	E02628	Genbank	2170856
Sequence 10771	E13330	Genbank	32521 <b>3</b> 5
Sequence 10772	J01415	Genbank	1944628
Sequence 10773	J03077	Genbank	183230
Sequence 10774	J03473	Genbank	337423
Sequence 10775	J03746	Genbank	183655
Sequence 10776	J04799	Genbank	190197
Sequence 10777	J04823	Genbank	1311703
Sequence 10778	J05016	Genbank	181507
Sequence 10779	J05480	Genbank	179807
Sequence 10780	L01057	Genbank	292462
Sequence 10781	L04483	Genbank	292444
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**Table 3-5** 

C 10780	1.05000	Genbank	388031
Sequence 10782	L05092 L05628	Genbank	1835658
Sequence 10783			
Sequence 10784	L08125	Genbank	306500
Sequence 10785	L09159	Genbank	307374
Sequence 10786	L11370	Genbank	387674
Sequence 10787	L12136	Genbank	181536
Sequence 10788	L12168	Genbank	178083
Sequence 10789	L19185	Genbank	440307
Sequence 10790	L20941	Genbank	507251
Sequence 10791	L28809	Genbank	454151
Sequence 10792	L29252	Genbank	808011
Sequence 10793	L31610	Genbank	1220360
Sequence 10794	L34160	Genbank	642091
Sequence 10795	L35681	Genbank	532031
	L38933	Genbank	1008841
Sequence 10796	L38951	Genbank	893287
Sequence 10797		Genbank	704415
Sequence 10798	L38995		
Sequence 10799	L39068	Genbank	994714
Sequence 10800	L39945	Genbank	703081
Sequence 10801	L40407	Genbank	703117
Sequence 10802	L41887	Genbank	950423
Sequence 10803	L47647	Genbank	1000861
Sequence 10804	L48984	Genbank	1066728
Sequence 10805	L78754	Genbank	1374832
Sequence 10806	M11147	Genbank	182513
Sequence 10807	M15887	Genbank	181960
Sequence 10808	M16342	Genbank	184266
Sequence 10809	M17886	Genbank	190233
Sequence 10810	M18930	Genbank	184371
Sequence 10811	M19989	Genbank	189719
Sequence 10812	M19997	Genbank	181968
Sequence 10813	M20496	Genbank	809235
Sequence 10814	M21142	Genbank	183402
Sequence 10815	M21154	Genbank	178517
Sequence 10816	M21575	Genbank	1311702
Sequence 10817	M24543	Genbank	341200
Sequence 10818	M24902	Genbank	189618
Sequence 10819	M25639	Genbank	188555
Sequence 10820	M26481	Genbank	619789
Sequence 10821	M26663	Genbank	618463
Sequence 10822	M27024	Genbank	341598
Sequence 10823	M29064	Genbank	337452
Sequence 10824	M29366	Genbank	181979
Sequence 10825	M29877	Genbank	178408
Sequence 10826	M31520	Genbank	337504
	M33146	Genbank	181070
Sequence 10827		Genbank	184381
Sequence 10828	M33616		180934
Sequence 10829	M34600	Genbank	
Sequence 10830	M36341	Genbank	178984
Sequence 10831	M37583	Genbank	184059
Sequence 10832	M58459	Genbank	337511
Sequence 10833	M58485	Genbank	180154
Sequence 10834	M60457	Genbank	181249
Sequence 10835	M60756	Genbank	184085
Sequence 10836	M64929	Genbank	190421
Sequence 10837	M69181	Genbank	641957

**Table 3-5** 

Sequence 10839         M76299         Gen           Sequence 10840         M81182         Gen           Sequence 10841         M81601         Gen           Sequence 10842         M83088         Gen           Sequence 10843         M93036         Gen           Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 187151 bank 188778 bank 190128 bank 339442 bank 189925 bank 182904 bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086 bank 493079
Sequence 10840         M81182         Gen           Sequence 10841         M81601         Gen           Sequence 10842         M83088         Gen           Sequence 10843         M93036         Gen           Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 188778 bank 190128 bank 339442 bank 189925 bank 182904 bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10841         M81601         Gen           Sequence 10842         M83088         Gen           Sequence 10843         M93036         Gen           Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 190128 bank 339442 bank 189925 bank 182904 bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10842         M83088         Gen           Sequence 10843         M93036         Gen           Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 339442 bank 189925 bank 182904 bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10842         M83088         Gen           Sequence 10843         M93036         Gen           Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 189925 bank 182904 bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10843         M93036         Gen           Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 182904 bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10844         M95775         Gen           Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 187180 bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10845         M96256         Gen           Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 182625 bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10846         M96982         Gen           Sequence 10847         S48220         Gen           Sequence 10848         S67861         Gen	bank 338262 bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10847         \$48220         Gen           Sequence 10848         \$67861         Gen	bank 257451 bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Sequence 10848 S67861 Gen	bank 239579 bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
_ •	bank 546602 bank 632789 bank 833998 bank 409404 bank 460086
Seculence MOGA S/M/90 (CAN)	bank 632789 bank 833998 bank 409404 bank 460086
0	bank 833998 bank 409404 bank 460086
A	bank 409404 bank 460086
0	bank 460086
0	
0 10071	
Sequence 10855 U09646 Gent	
Sequence 10856 U09813 Gent	
Sequence 10857 U12404 Gent	
Sequence 10858 U12465 Gent	
0	
Sequence 10859 U14650 Gent Sequence 10860 U14970 Gent	· · · · · · · · · · · · · · · · · · ·
Sequence 10861 U15008 Gent	
0	
Sequence 10862 U16797 Gent Sequence 10863 U17899 Gent	
Sequence 10864 U19144 Gent	
Sequence 10865 U21090 Gent	
Sequence 10866 U23803 Gent	
0	
Sequence 10867 U25182 Gent Sequence 10868 U28369 Gent	
Sequence 10869 U30826 Gent	
Sequence 10870 U32944 Gent	
Sequence 10871 U36336 Gent	
Sequence 10872 U38178 Gent	
Sequence 10873 U44754 Gent	
0	
Sequence 10874 U47742 Genb Sequence 10875 U49245 Genb	
0	
Sequence 10876 U49973 Genb Sequence 10877 U50939 Genb	
Sequence 10878 U52111 Genb	
Sequence 10879 U54993 Genb	
Sequence 10880 U55937 Genb	
Sequence 10881 U57847 Genb	
Sequence 10882 U58522 Genb	
Sequence 10883 U60644 Genb	
40000	
C	
10000	
0	
0	
10000	
0	
Sequence 10893 U80017 Genba	ank 1737211

**Table 3-5** 

Sequence 10894	U83463	Genbank	1916849
Sequence 10895	U87460	Genbank	2076881
Sequence 10896	U90441	Genbank	2439984
Sequence 10897	U90915	Genbank	1913895
Sequence 10898	U91323	Genbank	3582311
Sequence 10899	U91328	Genbank	2088550
Sequence 10900	U97670	Genbank	3264860
Sequence 10901	V00478	Genbank	28244
Sequence 10902	X00351	Genbank	28251
Sequence 10903	X03083	Genbank	34309
Sequence 10904	X04588	Genbank	37423
Sequence 10905	X05332	Genbank	35740
Sequence 10906	X13238	Genbank	1200056
Sequence 10907	X15187	Genbank	37260
Sequence 10908	X15341	Genbank	1197215
Sequence 10909	X15822	Genbank	30146
Sequence 10910	X51525	Genbank	30170
Sequence 10911	X53280	Genbank	29504
Sequence 10912	X53586	Genbank	33943
Sequence 10913	X56932	Genbank	23690
Sequence 10914	X58288	Genbank	32455
Sequence 10915	X60221	Genbank	509290
Sequence 10916	X63432	Genbank	28335
Sequence 10917	X63657	Genbank	296185
Sequence 10918	X64707	Genbank	29382
Sequence 10919	X65923	Genbank	31302
Sequence 10920	X67247	Genbank	36149
Sequence 10921	X67951	Genbank	287640
Sequence 10922	X74801	Genbank	671526
Sequence 10923	X74968	Genbank	439659
Sequence 10924	X75821	Genbank	509780
Sequence 10925	X75861	Genbank	456258
Sequence 10926	X78136	Genbank	460772
Sequence 10927	X78137	Genbank	460770
Sequence 10928	X78669	Genbank	469884
Sequence 10929	X80199	Genbank	2385366
Sequence 10930	X81109	Genbank	535057
Sequence 10931	X84694	Genbank	872117
Sequence 10932	X84908	Genbank	1502344
Sequence 10933	X86779	Genbank	1006658
Sequence 10934	X87176	Genbank	1050516
Sequence 10935	X87949	Genbank	1143491
Sequence 10936	X91790	Genbank	2511667
Sequence 10937	X97544	Genbank	1770563
Sequence 10938	X98356	Genbank	1648869
Sequence 10939	X99585	Genbank	1770518
Sequence 10940	Y00052	Genbank	30308
Sequence 10941	Y00282	Genbank	36048
Sequence 10942	Y00345	Genbank Genbank	35569 34266
Sequence 10943	Y00815		2181877
Sequence 10944 Sequence 10945	Y13620 Y15286	Genbank Genbank	2584788
	Z11692	Genbank Genbank	2564766 31107
Sequence 10946 Sequence 10947	Z11692 Z14000	Genbank Genbank	296063
Sequence 10947 Sequence 10948	Z14000 Z24724	Genbank Genbank	505034
Sequence 10949	Z36832	Genbank	533946
Cadneline 10343	سامان	Genuank	300340

**Table 3-5** 

Sequence 10950	Z37986	Genbank	780 <b>262</b>
Sequence 10951	Z37994	Genbank	556810
Sequence 10952	Z49250	Genbank	807889
Sequence 10953	Z63138	Genbank	1035516
Sequence 10954	Z64102	Genbank	1036480
Sequence 10955	Z68284	Genbank	1130698
Sequence 10956	Z70702	Genbank	1332492
Sequence 10957	Z82194	Genbank	2582744
Sequence 10958	Z83745	Genbank	1754650
Sequence 10959	Z83844	Genbank	4467 <b>20</b> 4
Sequence 10960	Z83856	Genbank	2995261
Sequence 10961	Z84722	Genbank	1817579
Sequence 10962	Z85999	Genbank	2326514
Sequence 10963	Z92544	Genbank	1869775
Sequence 10964	Z93016	Genbank	9650703
Sequence 10965	Z93242	Genbank	3164071
Sequence 10966	Z95114	Genbank	
Sequence 10967	Z95704	Genbank	5101742
Sequence 10968	Z97053		2121307
Sequence 10969	Z97056	Genbank	9650676
Sequence 10970	Z97183	Genbank	2832593
Sequence 10970	Z97630	Genbank	2961406
Sequence 10972	Z98749	Genbank	4582128
Sequence 10973	Z98882	Genbank	4775611
Sequence 10973		Genbank	4490865
Sequence 10975	Z98884 AA001709	Genbank	5304861
Sequence 10976	AA004303	EST	1445523
Sequence 10977	AA004303 AA007269	EST	1447948
Sequence 10978	AA007209 AA010173	EST	1463303
Sequence 10979	AA010173	EST	1471349
Sequence 10979		EST	1471260
Sequence 10980	AA010519 AA011539	EST	1471545
Sequence 10982		EST	1472565
Sequence 10983	AA015585 AA017170	EST	1476651
Sequence 10984	AA018892	EST	1479335
Sequence 10985	AA021623	EST	1482283
Sequence 10986	AA021023 AA025361	EST	1485284
Sequence 10987	AA026455	EST	1489375
Sequence 10988	AA026758	EST	1492355
Sequence 10989	AA027906	EST	1492556
Sequence 10990	AA031317	EST	1493993
Sequence 10991	AA035377	EST	1501272
Sequence 10992	AA037143	EST EST	1507101
Sequence 10993	AA039734		1512251
Sequence 10994	AA039957	EST	1516236
Sequence 10995	AA040714	EST	1516 <b>261</b>
Sequence 10996	AA044237	EST	1516992
Sequence 10997	AA044237 AA044575	EST	1522113
Sequence 10998	AA044703	EST	1522808
Sequence 10999		EST	1522971
Sequence 11000	AA045239	EST	1523443
Sequence 11000	AA045616	EST	1525379
Sequence 11002	AA047046	EST	1524944
Sequence 11002	AA054272	EST	1545197
Sequence 11003	AA057541	EST	1550246
Sequence 11004 Sequence 11005	AA058950	EST	1551758
Ocducinos i 1000	AA062994	EST	155 <b>7655</b>

**Table 3-5** 

Sequence 11006	AA063190	EST	1556924
Sequence 11007	AA063198	EST	1556796
Sequence 11008	AA070463	EST	1577823
Sequence 11009	AA070785	EST	1578347
Sequence 11010	AA074614	EST	1614492
Sequence 11011	AA075478	EST	1615543
Sequence 11012	AA076016	EST	1615903
Sequence 11013	AA083482	EST	1625543
Sequence 11014	AA085007	EST	1627125
Sequence 11015	AA085810	EST	1629342
Sequence 11016	AA088344	EST	1633856
Sequence 11017	AA089630	EST	1636122
Sequence 11018	AA093542	EST	1639127
Sequence 11019	AA095108	EST	1640693
Sequence 11020	AA098799	EST	1648802
Sequence 11021	AA099464	EST	1645408
Sequence 11022	AA099750	EST	1645861
Sequence 11023	AA100034	EST	1647326
Sequence 11024	AA101634	EST	1648570
Sequence 11024	AA101663	EST	1648744
Sequence 11025	AA102383	EST	1647229
Sequence 11026	AA102507	EST	1647621
Sequence 11027	AA113782	EST	1667703
Sequence 11028	AA114830	EST	1669952
Sequence 11029	AA115175	EST	1670372
Sequence 11030	AA115175 AA115455	EST	1670257
Sequence 11031	AA115455 AA115636	EST	1670237
Sequence 11032	AA115030 AA115738	EST	1670751
Sequence 11033		EST	1678078
Sequence 11034	AA122059	EST	1678318
Sequence 11035	AA122097	EST	1687948
Sequence 11036	AA126927		1692993
Sequence 11037	AA131505	EST EST	1693072
Sequence 11038	AA131583	EST	1690351
Sequence 11039	AA133311		1696705
Sequence 11040	AA135694	EST EST	1697294
Sequence 11041	AA136084		1697545
Sequence 11042	AA136337	EST	1697007
Sequence 11043	AA137210	EST	1712764
Sequence 11044	AA143393	EST	1716058
Sequence 11045	AA146684	EST	1720380
Sequence 11046	AA149579	EST	1720360
Sequence 11047	AA150361	EST	
Sequence 11048	AA151438	EST	1719775
Sequence 11049	AA151565	EST	1720183
Sequence 11050	AA151816	EST	1720511
Sequence 11051	AA152016	EST	1720854
Sequence 11052	AA156951	EST	1728584
Sequence 11053	AA157731	EST	1729356
Sequence 11054	AA160577	EST	1735945
Sequence 11055	AA161189	EST	1735443
Sequence 11056	AA161337	EST	1735652
Sequence 11057	AA165446	EST	1741462
Sequence 11058	AA169597	EST	1747985
Sequence 11059	AA169657	EST	1748026
Sequence 11060	AA172193	EST	1751270
Sequence 11061	AA173705	EST	1753883
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<u>Table 3-5</u>

Sequence 11062	AA174161	EST	1754441
Sequence 11063	AA176485	EST	1757618
Sequence 11064	AA177128	EST	1758286
Sequence 11065	AA178986	EST	1760538
Sequence 11066	AA180264	EST	1761531
Sequence 11067	AA186328	EST	1774446
Sequence 11068	AA187076	EST	1775245
Sequence 11069	AA188297	EST	1775383
Sequence 11070	AA190560	EST	1779535
Sequence 11071	AA193327	EST	1782738
Sequence 11072	AA193455	EST	1782846
Sequence 11073	AA203319	EST	1799028
Sequence 11074	AA206446	EST	1801826
Sequence 11075	AA207077	EST	1802642
Sequence 11076	AA209329	EST	1807273
Sequence 11077	AA211900	EST	1810537
Sequence 11078	AA223613	EST	1844155
Sequence 11079	AA225958	EST	1847285
Sequence 11080	AA226672	EST	1847988
Sequence 11081	AA232991	EST	1856115
Sequence 11082	AA233294	EST	1856287
Sequence 11083	AA234637	EST	1859306
Sequence 11084	AA235298	EST	1859735
Sequence 11085	AA243646	EST	1874456
Sequence 11086	AA248470	EST	1879307
Sequence 11087	AA248684	EST	1879900
Sequence 11088	AA250936	EST	1886035
Sequence 11089	AA251061	EST	1886024
Sequence 11090	AA251263	EST	1886226
Sequence 11091	AA251692	EST	1886655
Sequence 11092	AA252143	EST	1887106
Sequence 11093	AA252608	EST	1887582
Sequence 11094	AA252943	EST	1882696
Sequence 11095 Sequence 11096	AA255735	EST	1892673
Sequence 11097	AA257016	EST	1891145
Sequence 11097	AA258732 AA262091	EST	1893893
Sequence 11099	AA262425	EST	1898235
Sequence 11100	AA262683	EST	1897921
Sequence 11101	AA278796	EST	1898095
Sequence 11102	AA282285	EST	1920117
Sequence 11103	AA285126	EST	1925201
Sequence 11104	AA286765	EST	1928107
Sequence 11105	AA293B22	EST EST	1933628
Sequence 11106	AA296237	EST	1941 <b>74</b> 5 1948581
Sequence 11107	AA296690	EST	
Sequence 11108	AA296B12	EST	1949182
Sequence 11109	AA296923	EST	1949146
Sequence 11110	AA304244	EST	1949255
Sequence 11111	AA305213	EST	1956597
Sequence 11112	AA305921	EST	1957540
Sequence 11113	AA306323	EST	1958412
Sequence 11114	AA306941	EST	1958671
Sequence 11115	AA307728	EST	1959271
Sequence 11116	AA308688	EST	1960056 1961018
Sequence 11117	AA309119	EST	1961444
•		LUI	1301444

Sequence 11118	AA309929	EST	1962484
Sequence 11119	AA310375	EST	1962704
Sequence 11120	AA311577	EST	1963980
Sequence 11121	AA313294	EST	1965623
Sequence 11122	AA314382	EST	1966731
Sequence 11123	AA314635	EST	1967114
Sequence 11124	AA315157	EST	1967486
Sequence 11125	AA315989	EST	1968318
Sequence 11126	AA316456	EST	1968785
Sequence 11127	AA316676	EST	1969004
Sequence 11128	AA325081	EST	1977579
Sequence 11129	AA325713	EST	1977956
Sequence 11130	AA329695	EST	1981937
Sequence 11131	AA331249	EST	1983730
Sequence 11132	AA340295	EST	1992760
Sequence 11133	AA340744	EST	1992983
Sequence 11134	AA343834	EST	1996133
Sequence 11135	AA345527	EST	1997837
Sequence 11136	AA355326	EST	2007708
Sequence 11137	AA364646	EST	2016963
Sequence 11138	AA365552	EST	2017902
Sequence 11139	AA371377	EST	2023695
Sequence 11140	AA379942	EST	2032282
Sequence 11141	AA382161	EST	2034479
Sequence 11142	AA384806	EST	2037123
Sequence 11143	AA393525	EST	2046493
Sequence 11144	AA400534	EST	2054467
Sequence 11145	AA400644	EST	2054515
Sequence 11146	AA402038	EST	2056038
Sequence 11147	AA402743	EST	2057061
Sequence 11148	AA402800	EST	2056553
Sequence 11149	AA404235	EST	2058977
Sequence 11150	AA404535	EST	2059268
Sequence 11151	AA406310	EST	2064294
Sequence 11152	AA410936	EST	2070190
Sequence 11153	AA411736	EST	2069397
Sequence 11154	AA417324	EST	2077423
Sequence 11155	AA417743	EST	2079544
Sequence 11156	AA418847	EST ·	2080666
Sequence 11157	AA420643	EST	2094559
Sequence 11158	AA420650	EST	2094538
Sequence 11159	AA424006	EST	2102976
Sequence 11160	AA424349	EST	2103336
Sequence 11161	AA425535	EST	2106293
Sequence 11162	AA426021	EST	2106545
Sequence 11163	AA427560	EST	2111428
Sequence 11164	AA428565	EST	2112562
Sequence 11165	AA430605	EST	2111178
Sequence 11166	AA432282	EST	2114670
Sequence 11167	AA436100	EST	2141014
Sequence 11168	AA437224	EST	2142138
Sequence 11169	AA442070	EST	2153948
Sequence 11170	AA443297	EST	2155972
Sequence 11171	AA443762	EST	2156437
Sequence 11172	AA446702	EST	2159367
Sequence 11173	AA446759	EST	2159424
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Sequence 11174	AA446859	EST	2159524
Sequence 11175	AA446942	EST	2159607
Sequence 11176	AA447406	EST	2161076
Sequence 11177	AA447871	EST	2161541
Sequence 11178	AA448125	EST	2161795
Sequence 11179	AA448292	EST	2161962
Sequence 11180	AA449087	EST	2163107
Sequence 11181	AA450129	EST	2163879
Sequence 11182	AA451779	EST	216544B
Sequence 11183	AA452120	EST	2165789
Sequence 11184	AA454989	EST	
Sequence 11185	AA455869	EST	2177765
Sequence 11186	AA456454	EST	2178645
Sequence 11187	AA460766	EST	2179030
Sequence 11188	AA464281	_	2185886
Sequence 11189	AA464409	EST	2189165
		EST	2189293,
Sequence 11190	AA464803	EST	2189687
Sequence 11191	AA468360	EST	2194894
Sequence 11192	AA476267	EST	2204478
Sequence 11193	AA478669	EST	2207303
Sequence 11194	AA483131	EST	2211380
Sequence 11195	AA483251	EST	2212064
Sequence 11196	AA483450	EST	2212263
Sequence 11197	AA484155	EST	2212968
Sequence 11198	AA484907	EST	2213974
Sequence 11199	AA485065	EST	2214284
Sequence 11200	AA485608	EST	2214827
Sequence 11201	AA487694	EST	2217858
Sequence 11202	AA489068	EST	2218670
Sequence 11203	AA489782	EST	2220666
Sequence 11204	AA491858	EST	2221420
Sequence 11205	AA492143	EST	2221705
Sequence 11206	AA492423	EST	2221985
Sequence 11207	AA493522	EST	2223363
Sequence 11208	AA493635	EST	2223476
Sequence 11209	AA502361	EST	2237328
Sequence 11210	AA502517	EST	2237484
Sequence 11211	AA503033	EST	2238000
Sequence 11212	AA503078	EST	2238045
Sequence 11213	AA504572	EST	2240732
Sequence 11214	AA504890	EST	2241050
Sequence 11215	AA508394	EST	2245897
Sequence 11216	AA508597	EST	2246100
Sequence 11217	AA513123	EST	2251535
Sequence 11218	AA516472	EST	2255996
Sequence 11219	AA523172	EST	2263884
Sequence 11220	AA524187	EST	2265115
Sequence 11221	AA525960	EST	2268029
Sequence 11222	AA526740	EST	2268809
Sequence 11223	AA527181	EST	2269250
Sequence 11224	AA527494	EST	2269563
Sequence 11225	AA527667	EST	2269736
Sequence 11226	AA528148	EST	2270217
Sequence 11227	AA531585	EST	2274291
Sequence 11228	AA532665	EST	2276919
Sequence 11229	AA532805	EST	2278381

## <u>Table 3-5</u>

Sequence 11230	AA533279	EST	2277375
Sequence 11231		EST	2279351
Sequence 11232		EST	2287971
Sequence 11233		EST	2321393
Sequence 11234		EST	2321637
Sequence 11235		EST	2321961
Sequence 11236		EST	2322022
Sequence 11237		EST	2327651
Sequence 11238		EST	2327668
Sequence 11239		EST	2328110
Sequence 11240		EST	2331397
Sequence 11241	AA563778	EST	2335417
Sequence 11242	AA564118	EST	2335757
Sequence 11243	AA565432	EST	2337071
Sequence 11244		EST	2343826
Sequence 11245		EST	2348236
Sequence 11246		EST	2357670
Sequence 11247		EST	2355816
Sequence 11248		EST	2360684
Sequence 11249		EST	2359612
Sequence 11250		EST	2359730
Sequence 11251		EST	2368531
Sequence 11252		EST	
Sequence 11253		EST	2401716
Sequence 11254			2409164
		EST EST	2410689
Sequence 11255		EST	2432382
Sequence 11256			2433361
Sequence 11257		EST	2433819
Sequence 11258		EST	2436564
Sequence 11259		EST	2445248
Sequence 11260		EST	2457418
Sequence 11261		EST	2457890
Sequence 11262		EST	2458296
Sequence 11263		EST	2458852
Sequence 11264		EST	2464828
Sequence 11265		EST	2464071
Sequence 11266		EST	2464082
Sequence 11267		EST	2466215
Sequence 11268		EST	2505448
Sequence 11269		EST	2539022
Sequence 11270		EST	2539029
Sequence 11271		EST	2539220
Sequence 11272		EST	2540225
Sequence 11273		EST	2540838
Sequence 11274	AA628721	EST	2541108
Sequence 11275	AA630616	EST	2553227
Sequence 11276	AA630981	EST	2553592
Sequence 11277	AA631024	EST	2553635
Sequence 11278	AA631684	EST	2554295
Sequence 11279		EST	2555948
Sequence 11280		EST	2557008
Sequence 11281		EST	2557301
Sequence 11282		EST	2557334
Sequence 11283		EST	2565971
Sequence 11284		EST ·	2584128
Sequence 11285		EST	2584481

# <u>Table 3-5</u>

Sequence 11286	AA653775	EST	2589929
Sequence 11287	AA654914	EST	2591068
Sequence 11288	AA658095	EST	2594249
Sequence 11289	AA659440	EST	2595594
Sequence 11290	AA682466	EST	2616557
			2616621
Sequence 11291	AA662530	EST	
Sequence 11292	AA663778	EST	2617769
Sequence 11293	AA666385	EST	2620998
Sequence 11294	AA669098	EST	2630597
Sequence 11295	AA677560	EST	2658082
Sequence 11296	AA678246	EST	2658768
Sequence 11297	AA679533	EST	2660055
Sequence 11298	AA701882	EST	2704995
Sequence 11299	AA702803	EST	2705916
Sequence 11300	AA702824	EST	2705937
Sequence 11301	AA705800	EST	2715718
Sequence 11302	AA707319	EST	2717237
Sequence 11303	AA707321	EST	2717239
Sequence 11304	AA723194	EST	2740971
Sequence 11305	AA724307	EST	2742014
•	AA732439	EST	2753046
Sequence 11306	AA741013	EST	2779605
Sequence 11307	AA741421	EST	2780013
Sequence 11308		EST	2783511
Sequence 11309	AA744747		2788004
Sequence 11310	AA748046	EST	
Sequence 11311	AA748621	EST	2788579
Sequence 11312	AA761266	EST	2810196
Sequence 11313	AA761352	EST	2810282
Sequence 11314	AA768205	EST	2819220
Sequence 11315	AA768918	EST	2820156
Sequence 11316	AA773040	EST	2824611
Sequence 11317	AA775624	EST	2834958
Sequence 11318	AA775681	EST	2835015
Sequence 11319	AA777245	EST	2836576
Sequence 11320	AA777969	EST	2837370
Sequence 11321	AA778560	EST	2837891
Sequence 11322	AA779371	EST	2838702
Sequence 11323	AA782583	EST	2841914
Sequence 11324	AA788913	EST	2849033
Sequence 11325	AA804448	EST	2875961
Sequence 11326	AA805260	EST	2874010
Sequence 11327	AA805691	EST	2874441
Sequence 11328	AA809388	EST	2878794
Sequence 11329	AA809587	EST	2878993
Sequence 11330	AA810838	EST	2880449
Sequence 11331	AA812263	EST	2881874
Sequence 11332	AA813882	EST	2883478
Sequence 11333	AA814716	EST	2884312
Sequence 11334	AA827816	EST	2900179
Sequence 11335	AA829667	EST	2902766
Sequence 11336	AA831192	EST	2904291
Sequence 11337	AA834216	EST	2907815
Sequence 11338	AA837128	EST	2912327
Sequence 11339	AA838535	EST	2913334
Sequence 11340	AA845554	EST	2933313
Sequence 11341	AA846156	EST	2932296

Sequence 11342	AA846339	EST	2932479
Sequence 11343	AA846735	EST	2932875
Sequence 11344	AA846815	EST	2932955
Sequence 11345	AA847840	EST	2934358
Sequence 11346	AA852960	EST	2939699
•			
Sequence 11347	AA856586 AA864573	EST	2944888
Sequence 11348		EST	2958886
Sequence 11349	AA865331	EST	2957607
Sequence 11350	AA865871	EST	2958147
Sequence 11351	AA868273	EST	2963718
Sequence 11352	AA868720	EST	2964165
Sequence 11353	AA868840	EST	2964285
Sequence 11354	AA872275	EST	2968453
Sequence 11355	AA878211	EST	2987176
Sequence 11356	AA878424	EST	2987389
Sequence 11357	AA889466	EST	3016345
Sequence 11358	AA902388	EST	3037511
Sequence 11359	AA902974	EST	3038097
Sequence 11360	AA903453	EST	3038576
Sequence 11361	AA905040	EST	3040163
Sequence 11362	AA922765	EST	3070074
Sequence 11363	AA929066	EST	3078045
Sequence 11364	AA935780	EST	3092937
Sequence 11365	AA935796	EST	3092953
Sequence 11366	AA939273	EST	3099186
Sequence 11367	AA946596	EST	3109991
Sequence 11368	AA961071	EST	3127625
Sequence 11369	AA962465	EST	3134629
Sequence 11370	AA973265	EST	3148445
Sequence 11371	AA975592	EST	3151384
Sequence 11372	AA977036	EST	3154482
Sequence 11373	AA977991	EST	3155437
Sequence 11374	AA989018	EST	3174589
Sequence 11375	AA993218	EST	3179763
Sequence 11376	AA993706	EST	3180251
Sequence 11377	AF109301	EST	6782694
Sequence 11378	Al002701	EST	3203115
Sequence 11379	Al002736	EST	3203150
Sequence 11380	AI016055	EST	3230391
Sequence 11381	Al017469	EST	3231805
Sequence 11382	Al018676	EST	3232474
Sequence 11383	Al025274	EST	3240887
Sequence 11384	AI027434	EST	3244950
Sequence 11385	Al028115		
	Al028439	EST	3245424
Sequence 11386	AI033712	EST	3245748
Sequence 11387		EST	3254665
Sequence 11388	A1034384	EST	3255337
Sequence 11389	A1034428	EST	3255381
Sequence 11390	A1039027	EST	3278221
Sequence 11391	Al039136	EST	3278330
Sequence 11392	AI039731	EST	3278925
Sequence 11393	Al040803	EST	3279997
Sequence 11394	Al051985	EST	3307976
Sequence 11395	Al052598	EST	3308589
Sequence 11396	AI053744	EST	3321531
Sequence 11397	AI056319	EST	3330185

**Table 3-5** 

Sequence 11398	AI056324	EST	3330190
Sequence 11399	AI056337	EST	3330203
Sequence 11400	AI056351	EST	3330217
Sequence 11401	AI064691	EST	6358963
Sequence 11402	AI065139	EST	6359411
Sequence 11403	AI066536	EST	3367238
Sequence 11404	AI073929	EST	3400573
Sequence 11405	AI073929 AI074143	EST	3400787
Sequence 11406	AI074143 AI076009	EST	3405187
•	AI079555		
Sequence 11407		EST	3415806
Sequence 11408	AI079831	EST	3416082
Sequence 11409	AI079885	EST	3416136
Sequence 11410	A1080266	EST	3416517
Sequence 11411	A1084035	EST	3422458
Sequence 11412	AI086937	EST	3425360
Sequence 11413	AI088270	EST	3427329
Sequence 11414	AI089408	EST	34 <b>28467</b>
Sequence 11415	AI089953	EST	34 <b>29012</b>
Sequence 11416	Al091326	EST	3430385
Sequence 11417	AI091580	EST	3430639
Sequence 11418	Al091864	EST	3430923
Sequence 11419	AI092638	EST	3431614
Sequence 11420	Al096462	EST	3445956
Sequence 11421	Al096678	EST	3446172
Sequence 11422	Al114651	EST	6359996
Sequence 11423	Al125541	EST	3594055
Sequence 11424	Al125669	EST	3594183
Sequence 11425	Al127461	EST	3595975
Sequence 11426	Al129721	EST	3598235
Sequence 11427	Al130978	EST	3600994
Sequence 11428	Al131385	EST	3601401
Sequence 11429	Al141972	EST	3649429
Sequence 11430	Al143723	EST	3665532
Sequence 11431	Al144216	EST	3666025
Sequence 11432	Al147846	EST	3675528
Sequence 11433	Al149616	EST	3678085
Sequence 11434	Al150878	EST	3679347
Sequence 11435	Al184622	EST	
Sequence 11436			3735260
	Al187296	EST	3737934
Sequence 11437	Al187413	EST	3738051
Sequence 11438	Al187993	EST	3739202
Sequence 11439	Al188768	EST	3739977
Sequence 11440	Al190574	EST	3741783
Sequence 11441	Al193729	EST	3744938
Sequence 11442	Al193879	EST	3745088
Sequence 11443	Al202702	EST	3755308
Sequence 11444	Al219886	EST	3802089
Sequence 11445	Al222521	EST	3804724
Sequence 11446	Al225043	EST	3807756
Sequence 11447	Al243691	EST	3839088
Sequence 11448	Al244187	EST	3839584
Sequence 11449	Al251103	EST	3847632
Sequence 11450	Al253330	EST	3850451
Sequence 11451	Al253436	EST	3850391
Sequence 11452	Al262640	EST	3870843
Sequence 11453	Al263331	EST	3871534
- · · · · · · · · · · · · · · · · · · ·			JJ. 100-7

**Table 3-5** 

Sequence 1	1454	AI267499	EST	3886666
Sequence 1		Al269682	EST	3888849
Sequence 1		Al270013	EST	3889180
Sequence 1		Al271339	EST	3890506
Sequence 1		Al271608	EST	3890775
Sequence 1		Al273048	EST	3895316
Sequence 1		Al275943	EST	3898217
Sequence 1		AI278640	EST	3916874 3917232
Sequence 1		Al278998	EST	
Sequence 1		AI280181	EST	3918414
Sequence 1		Al281762	EST	3919995
Sequence 1		Al282246	EST	3920479
Sequence 1		A1282856	EST	3921089
Sequence 1		Al284545	EST	3922778
Sequence 1	1468	AI289108	EST	3932372
Sequence 1	1469	Al290847	EST	3933621
Sequence 1	1470	Al301608	EST	3960954
Sequence 1		Al306464	EST	3989535
Sequence 1		Al308012	EST	4002647
Sequence 1		Al312384	EST	4017989
Sequence 1		Al339601	EST	4076528
Sequence 1		Al347153	EST	4084359
Sequence 1		Al352526	EST	4089732
Sequence 1		Al356383	EST	4108004
Sequence 1		Al358021	EST	4109642
Sequence 1		Al358557	EST	4110178
Sequence 1		Al359812	EST	4111433
Sequence 1		Al362921	EST	4114542
Sequence 1		Al366222	EST	4125911
Sequence 1		Al393262	EST	4222809
Sequence 1		Al393754	EST	4223301
Sequence 1		Al400359	EST	4243446
Sequence 1		Al417686	EST	4261190
Sequence 1		Al420543	EST	4266474
Sequence 1		AI436691	EST	4283721
Sequence 1		Al439350	EST	4303707
Sequence 1		Al468004	EST	4330094
		AI469347	EST	4331437
Sequence 1		A1469505	EST	4331595
Sequence 1		A1469600		4331690
Sequence 1		AI472201	EST EST	4334291
Sequence 1				
Sequence 1		AI478202	EST	4371428
Sequence 1		Al479751	EST	4372919
Sequence 1		AI480256	EST	4373424
Sequence 1		A1497917	EST	4389899
Sequence 1		A1499665	EST	4391647
Sequence 1		AI521246	EST	4435381
Sequence 1		AI525607	EST	4439742
Sequence 1		AI539780	EST	4453915
Sequence 1		AI559444	EST	4509649
Sequence 1		AI559531	EST	4509736
Sequence 1		A1565555	EST	4524012
Sequence 1		AI570526	EST	4533900
Sequence 1		AI582754	EST	4568651
Sequence 1		A1608932	EST	4618099
Sequence 1	150 <del>9</del>	AI608936	EST	4618103

Sequence 11510	Al621173	EST	4630299
Sequence 11511	Al623863	EST	4648794
Sequence 11512	Al625152	EST	4650083
Sequence 11513		EST	4650750
Sequence 11514		EST	
Sequence 11515		EST	4683326
Sequence 11516	Al635397		4684285
Sequence 11517	Al635756	EST	4686727
Sequence 11518		EST	4687086
Sequence 11519	AI657111	EST	4741090
	Al683799	EST	4893981
Sequence 11520	AI684760	EST	4896054
Sequence 11521	Al699114	EST	4987014
Sequence 11522	AI700656	EST	4988556
Sequence 11523	AI734838	EST	5056362
Sequence 11524	Al741189	EST	5109477
Sequence 11525	Al743974	EST	5112262
Sequence 11526	Al744342	EST	5112630
Sequence 11527	Al751021	EST	5129373
Sequence 11528	Al751175	EST	5129439
Sequence 11529	Al762495	EST	5178162
Sequence 11530	AI763344	EST	5179011
Sequence 11531	Al765690	EST	5232199
Sequence 11532	A1799995	EST	5365467
Sequence 11533	Al802542	EST	5368014
Sequence 11534	AI804082	EST	
Sequence 11535	AI808503	EST	5369554
Sequence 11536	Al813654	EST	5395069
Sequence 11537	AI815499		5424869
Sequence 11538	AI858089	EST	5431045
Sequence 11539	AI866603	EST	5511705
Sequence 11540	AI866803	EST	5530710
Sequence 11541	A1867464	EST	5530834
Sequence 11542		EST	5540480
Sequence 11543	AI879010	EST	5553059
Sequence 11544	AI887361	EST	5592525
	Al903990	EST	6494286
Sequence 11545	Al905002	EST	6495389
Sequence 11546	Al929357	EST	5665321
Sequence 11547	Al929483	EST	5665447
Sequence 11548	Al948496	EST	5740806
Sequence 11549	Al961342	EST	5754055
Sequence 11550	Al983520	EST	5810739
Sequence 11551	AJ133829	EST	4727122
Sequence 11552	AL036039	EST	5405665
Sequence 11553	AL036415	EST	5405971
Sequence 11554	AL037828	EST	5407161
Sequence 11555	AL039773	EST	5408788
Sequence 11556	AL042955	EST	5422386
Sequence 11557	AL043992	EST	5432220
Sequence 11558	AL048634	EST	4727722
Sequence 11559	AL119245	EST	
Sequence 11560	AL120824	EST	5925144 5026925
Sequence 11561	AL134867	EST	5926825
Sequence 11562	AL135047		6603054
Sequence 11563	AL138273	EST	6603234
Sequence 11564	AL138451	EST	6854954
Sequence 11565	AW003648	EST	6855132
- 34401100 1 1000	AVV 003048	EST	5850564

			5050007
Sequence 11566	AW008029	EST	5856807
Sequence 11567	AW008048	EST	5856826
Sequence 11568	AW008782	EST	5857560
Sequence 11569	AW021322	EST	5874782
Sequence 11570	AW021588	EST	5875118
Sequence 11571	AW022159	EST	5875689
Sequence 11572	AW043631	EST	5904160
Sequence 11573	AW044310	EST	5904839
Sequence 11573	AW069285	EST	6024283
Sequence 11574	AW069315	EST	6024313
Sequence 11575	AW073580	EST	6028578
Sequence 11576	AW082238	EST	6037390
Sequence 11577	AW083099	EST	6038251
Sequence 11578	AW085581	EST	6040733
Sequence 11579	AW 083361 AW 087200	EST	6043005
Sequence 11580		EST	6140775
Sequence 11581	AW136642	EST	6196586
Sequence 11582	AW148690		6229259
Sequence 11583	AW 157858	EST	6299739
Sequence 11584	AW 160706	EST	6300361
Sequence 11585	AW161328	EST	6301976
Sequence 11586	AW162943	EST	•••
Sequence 11587	AW167129	EST	6398654
Sequence 11588	AW168050	EST	6399575
Sequence 11589	AW168184	EST	6399709
Sequence 11590	AW168400	EST	6399925
Sequence 11591	AW 177050	EST	6443087
Sequence 11592	AW 192594	EST	6471293
Sequence 11593	AW 205906	EST	6505380
Sequence 11594	AW247236	EST	6590229
Sequence 11595	AW250104	EST	6593097
Sequence 11596	AW294606	EST	6701242
Sequence 11597	AW294626	EST	6701262
Sequence 11598	AW 297943	EST	6704579
Sequence 11599	AW302841	EST	6712521
Sequence 11600	AW337419	EST	6834 <b>0</b> 45
Sequence 11601	AW363949	EST	6868599
Sequence 11602	AW363975	EST	6868625
Sequence 11603	AW365134	EST	6869784
Sequence 11604	AW365960	EST	6870610
Sequence 11605	AW368992	EST	6873642
Sequence 11606	AW376614	EST	6881275
Sequence 11607	AW380617	EST	6885276
Sequence 11608	AW380631	EST	6885290
Sequence 11609	AW381959	EST	6886618
Sequence 11610	AW386920	EST	6891579
Sequence 11611	AW401953	EST	6920639
Sequence 11612	AW402499	EST	6921188
Sequence 11613	AW 450642	EST	6991418
Sequence 11013	AW452921	EST	6993697
Sequence 11614	AW 499890	EST	7111988
Sequence 11615	AW 503119	EST	7118224
Sequence 11616	AW518064	EST	N/A
Sequence 11617		EST	7238326
Sequence 11618	AW573593	EST	7246279
Sequence 11619	AW574740	EST	7246633
Sequence 11620	AW575094	EST	7251834
Sequence 11621	AW576694	E91	7231034

Sequence 11622	AW579819	EST	7254868
Sequence 11623	D44813	EST	1572288
Sequence 11624	D55269	EST	957166
Sequence 11625	D80421	EST	
Sequence 11626	F07786		1178298
Sequence 11627	F20959	EST	677286
Sequence 11628		EST	2060135
	F22113	EST	2061289
Sequence 11629	H01421	EST	864354
Sequence 11630	H11779	EST	876599
Sequence 11631	H16460	EST	881280
Sequence 11632	H16546	EST	882771
Sequence 11633	H18560	EST	884800
Sequence 11634	H22094	EST	890789
Sequence 11635	H40387	EST	916439
Sequence 11636	H54240	EST	994387
Sequence 11637	H68450	EST	1027190
Sequence 11638	H90819	EST	1081249
Sequence 11639	H94986	EST	
Sequence 11640	L44419	EST	1102619
Sequence 11641	N20934	EST	1048740
Sequence 11642	N21624		1126104
Sequence 11643	N23545	EST	1126794
Sequence 11644	N26941	EST	1137695
Sequence 11645	N28426	EST	1141289
Sequence 11646	N31770	EST	1146662
Sequence 11647	N40973	EST	1152169
Sequence 11648	N44346	EST	1164571
Sequence 11649		EST	1182865
Sequence 11650	N54935	EST	1196255
Sequence 11651	N71715	EST	1228427
Sequence 11652	N80408	EST	1243109
Sequence 11653	R17938	EST	771548
	R82925	EST	927634
Sequence 11654	R93415	EST	967581
Sequence 11655	R93745	EST	967911
Sequence 11656	T30702	EST	612800
Sequence 11657	T48489	EST	642689
Sequence 11658	T77803	EST	695006
Sequence 11659	T89533	EST	718046
Sequence 11660	W01580	EST	1273634
Sequence 11661	W01785	EST	1273784
Sequence 11662	W02490	EST	1274488
Sequence 11663	W33011	EST	1315016
Sequence 11664	W79178	EST	1390613
Sequence 11665	V60015	NUCPATENT	N/A
Sequence 11666	V86392	NUCPATENT	N/A
Sequence 11667	X40046	NUCPATENT	N/A
Sequence 11668	X83330	NUCPATENT	949933
Sequence 11669	X87396	NUCPATENT	1304366
Sequence 11670	X97973	NUCPATENT	1360128
Sequence 11671	X98409	NUCPATENT	1403131
Sequence 11672	Z33373	NUCPATENT	534033
Sequence 11673	Z33622	NUCPATENT	608977
Sequence 11674	Z33656	NUCPATENT	488829
Sequence 11675	Z39728	NUCPATENT	565523
Sequence 11676	AC02994	PREPATNUC	N/A
Sequence 11677	AC31122	PREPATNUC	N/A
			11/1

**Table 3-5** 

Sequence 11678	AC34163	PREPATNUC	N/A
Sequence 11679	AC35230	PREPATNUC	N/A
Sequence 11680	AC36946	PREPATNUC	N/A
Sequence 11681	A02759	Genbank	345130
Sequence 11682	A16794	Genbank	512417
Sequence 11683	A17003	Genbank	512419
Sequence 11684	A20255	Genbank	1247401
Sequence 11685	A36460	Genbank	2293778
Sequence 11686	AB000584	Genbank	1813326
Sequence 11687	AB001535	Genbank	3928755
Sequence 11688	AB002308	Genbank	6634012
Sequence 11689	AB002321	Genbank	2224586 2224590
Sequence 11690	AB002323	Genbank	2224590 2224718
Sequence 11691	AB002387	Genbank	2280487
Sequence 11692	AB002390	Genbank	1944124
Sequence 11693	AB002533	Genbank	1945608
Sequence 11694	AB003102	Genbank	3702678
Sequence 11695	AB003151	Genbank	2102636
Sequence 11696	AB003698	Genbank	2627128
Sequence 11697	AB003730	Genbank	3077761
Sequence 11698	AB007166	Genbank Genbank	2887418
Sequence 11699	AB007882	Genbank Genbank	2887428
Sequence 11700	AB007887		6683704
Sequence 11701	AB007916	Genbank Genbank	3413857
Sequence 11702	AB007917	Genbank	6634036
Sequence 11703	AB007922	Genbank	3413899
Sequence 11704	AB007938	Genbank	2967451
Sequence 11705	AB010882 AB011004	Genbank	3273315
Sequence 11706	AB011004 AB011083	Genbank	3043545
Sequence 11707	AB011111	Genbank	3043601
Sequence 11708	AB011139	Genbank	3043657
Sequence 11709	AB011420	Genbank	3834353
Sequence 11710	AB014077	Genbank	5672586
Sequence 11711 Sequence 11712	AB014554	Genbank	3327121
Sequence 11713	AB015639	Genbank	5821139
Sequence 11713	AB017004	Genbank	4239949
Sequence 11715	AB017103	Genbank	3493334
Sequence 11716	AB017335	Genbank	3582440
Sequence 11717	AB017708	Genbank	3650434
Sequence 11718	AB018260	Genbank	3882154
Sequence 11719	AB018265	Genbank	3882164
Sequence 11720	AB018267	Genbank	3882168
Sequence 11721	AB018304	Genbank	3882242
Sequence 11722	AB018330	Genbank	3882294
Sequence 11723	AB018344	Genbank	3882322
Sequence 11724	AB019524	Genbank	4519939
Sequence 11725	AB019568	Genbank	3885371
Sequence 11726	AB020662	Genbank	4240198
Sequence 11727	AB020679	Genbank	4240232
Sequence 11728	AB020684	Genbank	4240242
Sequence 11729	AB020686	Genbank	4240246
Sequence 11730	AB020692	Genbank	4240258
Sequence 11731	AB020697	Genbank	4240268
Sequence 11732	AB020715	Genbank	4240304
Sequence 11733	AB020858	Genbank	4003378
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Sequence 11734		Genbank	4003381
Sequence 11735		Genbank	4038732
Sequence 11736	AB021467	Genbank	4586518
Sequence 11737	AB021663	Genbank	4996450
Sequence 11738	AB022435	Genbank	4996607
Sequence 11739	AB022537	Genbank	6942217
Sequence 11740	AB023050	Genbank	
Sequence 11741	AB023691	Genbank	5672605
Sequence 11742	AB023699		6519222
Sequence 11743	AB024334	Genbank	6519298
Sequence 11744	AB026894	Genbank	6016837
Sequence 11745		Genbank	5821423
Sequence 11745	AB027466	Genbank	6172220
	AB028859	Genbank	6567165
Sequence 11747	AB029003	Genbank	5689496
Sequence 11748	AB029347	Genbank	5921132
Sequence 11749	AB029551	Genbank	6714542
Sequence 11750	AB032417	Genbank	6277265
Sequence 11751	AB032969	Genbank	6329965
Sequence 11752	AB032975	Genbank	6330044
Sequence 11753	AB033070	Genbank	6330818
Sequence 11754	AB033076	Genbank	6330860
Sequence 11755	AB033079	Genbank	6382025
Sequence 11756	AB033082	Genbank	6330910
Sequence 11757	AB033117	Genbank	6331435
Sequence 11758	AB034205	Genbank	6899845
Sequence 11759	AB037744	Genbank	7243026
Sequence 11760	AB037757	Genbank	7243052
Sequence 11761	AB037763	Genbank	7243064
Sequence 11762	AB037767	Genbank	7243064 7243072
Sequence 11763	AB037807	Genbank	7243072
Sequence 11764	AB037849	Genbank	7243132
Sequence 11765	AC000040	Genbank	
Sequence 11766	AC000083	Genbank	7923866 3810636
Sequence 11767	AC000093	Genbank	4678240
Sequence 11768	AC000120	Genbank	
Sequence 11769	AC000353	Genbank	1809224
Sequence 11770	AC000378	Genbank	6970735
Sequence 11771	AC000394	Genbank	2270906
Sequence 11772	AC001226	Genbank	2133911
Sequence 11773	AC002039	Genbank	2133862
Sequence 11774	AC002040	Genbank	2342716
Sequence 11775	AC002064	Genbank	2347081
Sequence 11776	AC002069	Genbank	2076723
Sequence 11777	AC002094	Genbank	2076718
Sequence 11778	AC002117	Genbank	2155224
Sequence 11779	AC002401	Genbank	2281075
Sequence 11780	AC002470		2599240
Sequence 11781	AC002470 AC002481	Genbank	6478944
Sequence 11782	AC002544	Genbank	2340092
Sequence 11783	AC002544 AC003041	Genbank	3337382
Sequence 11784	AC003041 AC003103	Genbank	3264572
Sequence 11785		Genbank	2842782
Sequence 11786	AC003109	Genbank	2961444
Sequence 11787	AC003111	Genbank	2636670
Sequence 11788	AC003682	Genbank	3264845
Sequence 11789	AC003688	Genbank	3789719
andreing 11/08	AC003964	Genbank	2981253

		Genbank	2772538
Sequence 11790	AC003989	Genbank	2772566
Sequence 11791	AC003999	Genbank	2781390
Sequence 11792	AC004006		2811101
Sequence 11793	AC004039	Genbank	3366580
Sequence 11794	AC004049	Genbank	4001528
Sequence 11795	AC004050	Genbank	3097872
Sequence 11796	AC004098	Genbank	
Sequence 11797	AC004131	Genbank	3342217
Sequence 11798	AC004142	Genbank	2880078
Sequence 11799	AC004148	Genbank	3482960
Sequence 11800	AC004150	Genbank	4585948
Sequence 11801	AC004156	Genbank	2896799
Sequence 11802	AC004221	Genbank	9665053
Sequence 11803	AC004228	Genbank	4263838
Sequence 11804	AC004258	Genbank	2924759
Sequence 11805	AC004263	Genbank	2935616
Sequence 11806	AC004381	Genbank	2982169
Sequence 11807	AC004408	Genbank	3873185
Sequence 11808	AC004467	Genbank	3132844
Sequence 11809	AC004472	Genbank	2984582
Sequence 11810	AC004474	Genbank	3097873
Sequence 11811	AC004509	Genbank	2996628
Sequence 11812	AC004520	Genbank	3004572
Sequence 11813	AC004525	Genbank	3219333
Sequence 11814	AC004527	Genbank	4760422
Sequence 11815	AC004530	Genbank	3108050
Sequence 11816	AC004531	Genbank	3337392
Sequence 11817	AC004552	Genbank	3273378
Sequence 11818	AC004583	Genbank	3293210
Sequence 11819	AC004584	Genbank	3417305
Sequence 11820	AC004643	Genbank	3097841
Sequence 11821	AC004656	Genbank	3253114
Sequence 11822	AC004660	Genbank	3108018
Sequence 11823	AC004686	Genbank	3688105
Sequence 11824	AC004742	Genbank	3152630
Sequence 11825	AC004749	Genbank	3157918
Sequence 11826	AC004754	Genbank	3165400
Sequence 11827	AC004771	Genbank	3668108
Sequence 11828	AC004776	Genbank	3169299
Sequence 11829	AC004810	Genbank	3264567
Sequence 11830	AC004812	Genbank	3970970
Sequence 11831	AC004821	Genbank	4753291
Sequence 11832	AC004832	Genbank	6624129
Sequence 11833	AC004839	Genbank	3983571
Sequence 11834	AC004841	Genbank	4454523
Sequence 11835	AC004846	Genbank	7243869
Sequence 11836	AC004854	Genbank	4827328
Sequence 11837	AC004858	Genbank	6624125
Sequence 11838	AC004859	Genbank	5091656
Sequence 11839	AC004883	Genbank	4263746
Sequence 11840	AC004890	Genbank	4508146
Sequence 11841	AC004904	Genbank	4156180
Sequence 11842	AC004912	Genbank	4156178
Sequence 11843	AC004943	Genbank	3924671
Sequence 11844	AC004950	Genbank	4753269
Sequence 11845	AC004985	Genbank	5708490
Sadnering 1 1943	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	- <del>-</del> - · · ·	

Sequence 1184	6 AC004990	Genbank	2004000
Sequence 1184:	7 AC004999	Genbank	3924668
Sequence 1184	B AC005034	Genbank	3970963
Sequence 11849	AC005037	Genbank	3947437
Sequence 11850	AC005041	Genbank	4827310
Sequence 11851	AC005042	Genbank	4508118
Sequence 11852	AC005084	Genbank	4156138
Sequence 11853		Genbank	3659503
Sequence 11854	AC005104	Genbank Genbank	5732140
Sequence 11855	AC005153	_	4218027
Sequence 11856	AC005210	Genbank Genbank	3242766
Sequence 11857	AC005257		6249673
Sequence 11858	AC005263	Genbank Genbank	3289994
Sequence 11859	AC005318		3289978
Sequence 11860	AC005324	Genbank	3885345
Sequence 11861	AC005329	Genbank	3366582
Sequence 11862	AC005332	Genbank	3342732
Sequence 11863	AC005369	Genbank	3659494
Sequence 11864	AC005382	Genbank	3367505
Sequence 11865	AC005488	Genbank	338 <b>6587</b>
Sequence 11866	AC005550	Genbank	5836194
Sequence 11867	AC005614	Genbank	3478668
Sequence 11868	AC005621	Genbank	3540175
Sequence 11869	AC005627	Genbank	3548788
Sequence 11870	AC005629	Genbank	7243867
Sequence 11871	AC005630	Genbank	7243877
Sequence 11872	AC005661	Genbank	4159882
Sequence 11873	AC005702	Genbank	3818348
Sequence 11874	AC005726	Genbank	3688100
Sequence 11875	AC005786	Genbank	3810672
Sequence 11876	AC005789	Genbank	3702287
Sequence 11877	AC005837	Genbank	3702281
Sequence 11878	AC005838	Genbank Genbank	3849820
Sequence 11879	AC005841	Genbank Genbank	3947427
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Sequence 11881	AC005971	Genbank	4165005
Sequence 11882	AC006008	Genbank	4314421
Sequence 11883	AC006014	Genbank	5091655
Sequence 11884	AC006017	Genbank	4699959
Sequence 11885	AC006040	Genbank	4508141
Sequence 11886	AC006057	Genbank	4314426
Sequence 11887	AC006115	Genbank	4731048
Sequence 11888	AC006137	Genbank	3962498
Sequence 11889	AC006160	Genbank	9887710
Sequence 11890	AC006213	Genbank	5701616
Sequence 11891	AC006230	Genbank	4160143
Sequence 11892	AC006255	Genbank	4966389
Sequence 11893	AC006273	Genbank	5306223
Sequence 11894	AC006299	Genbank	4096060
Sequence 11895	AC006475	Genbank	4106997
Sequence 11896	AC006477	Genbank	4753283
Sequence 11897	AC006480	Genbank	7243871
Sequence 11898	AC006512	Genbank	6289252
Sequence 11899	AC006566	Genbank	4926863
Sequence 11900	AC006581	Genbank Genbank	4309780
Sequence 11901	AC007032	Genbank	4914350
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Table 3-5

		<b>a</b> 1 -1-	E709471
Sequence 11902	AC007041	Genbank	5708471 4885691
Sequence 11903	AC007055	Genbank	4508098
Sequence 11904	AC007066	Genbank	4416547
Sequence 11905	AC007073	Genbank	4731066
Sequence 11906	AC007172	Genbank	4558643
Sequence 11907	AC007191	Genbank	6715703
Sequence 11908	AC007225	Genbank	5649375
Sequence 11909	AC007253	Genbank	5708473
Sequence 11910	AC007384	Genbank	4689442
Sequence 11911	AC007406	Genbank	5263309
Sequence 11912	AC007528	Genbank	4914348
Sequence 11913	AC007537	Genbank Genbank	4982536
Sequence 11914	AC007541		4982534
Sequence 11915	AC007564	Genbank Genbank	4835751
Sequence 11916	AC007565	Genbank	5732177
Sequence 11917	AC007736	Genbank	5306288
Sequence 11918	AC007938	Genbank	7007656
Sequence 11919	AC007999	Genbank	6015188
Sequence 11920	AC008044	Genbank	6970743
Sequence 11921	AC008101	Genbank	9797815
Sequence 11922	AC008102	Genbank	5931373
Sequence 11923	AC008122	Genbank	6006046
Sequence 11924	AC008123	Genbank	7243890
Sequence 11925	AC008269	Genbank	6850299
Sequence 11926	AC008498 AC008893	Genbank	7259691
Sequence 11927	AC009330	Genbank	6553960
Sequence 11928	AC010206	Genbank	6539154
Sequence 11929	AC010200 AC010582	Genbank	6721135
Sequence 11930	AC010302 AC011331	Genbank	9929686
Sequence 11931	AC012087	Genbank	7114465
Sequence 11932	AC020663	Genbank	6682593
Sequence 11933	AC021049	Genbank	7248920
Sequence 11934	AF001549	Genbank	3355302
Sequence 11935	AF001977	Genbank	2914738
Sequence 11936	AF002668	Genbank	2232173
Sequence 11937 Sequence 11938	AF003837	Genbank	2228792
Sequence 11939	AF004162	Genbank	3046385
Sequence 11940	AF004339	Genbank	2245559
Sequence 11941	AF004341	Genbank	2245563
Sequence 11942	AF004342	Genbank	2245565
Sequence 11943	AF005896	Genbank	2209244
Sequence 11944	AF006083	Genbank	2282031
Sequence 11945	AF007544	Genbank	2970122
Sequence 11946	AF007835	Genbank	4102034
Sequence 11947	AF008303	Genbank	2580619
Sequence 11948	AF011889	Genbank	2335186
Sequence 11949	AF012023	Genbank	2305237
Sequence 11950	AF012072	Genbank	9967556
Sequence 11951	AF012126	Genbank	2832227
Sequence 11952	AF015812	Genbank	2599359
Sequence 11953	AF016709	Genbank	2731560
Sequence 11954	AF019413	Genbank	2347130
Sequence 11955	AF020057	Genbank	3150436
Sequence 11956	AF020351	Genbank	2655052
Sequence 11957	AF020352	Genbank	2655054
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Sequence 11958		Genbank	2460317
Sequence 11959		Genbank	2558890
Sequence 11960	AF026292	Genbank	2559009
Sequence 11961	AF026816	Genbank	2583078
Sequence 11962	AF028832	Genbank	3287488
Sequence 11963	AF030186	Genbank	3831546
Sequence 11964	AF032387	Genbank	
Sequence 11965	AF033095	Genbank	2641556
Sequence 11966	AF035840	Genbank	2645728
Sequence 11967	AF036715	Genbank	3800739
Sequence 11968	AF038178	Genbank	4007385
Sequence 11969	AF038200	Genbank	2795898
Sequence 11970	AF038554	Genbank	2795921
Sequence 11971	AF038633	Genbank	2895558
Sequence 11972	AF038955	_	3252876
Sequence 11973	AF041483	Genbank	3329379
Sequence 11974	AF042166	Genbank	3493528
Sequence 11975	AF042836	Genbank	3298596
Sequence 11976	AF044957	Genbank	3776241
Sequence 11977		Genbank	4164445
Sequence 11978	AF044958 AF047185	Genbank	4164447
Sequence 11979	AF047185 AF047440	Genbank	2909861
Sequence 11980	AF047440 AF047472	Genbank	3335135
Sequence 11981		Genbank	2921872
Sequence 11982	AF049105	Genbank	2984656
Sequence 11983	AF051316	Genbank	6580759
Sequence 11984	AF051894	Genbank	3095110
Sequence 11985	AF052129	Genbank	3360438
Sequence 11986	AF052642	Genbank	2981453
Sequence 11987	AF053356	Genbank	3135305
Sequence 11988	AF054984	Genbank	3005692
Sequence 11989	AF054990	Genbank	3005703
Sequence 11990	AF055473	Genbank	nil
Sequence 11991	AF055474	Genbank	nil
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Sequence 11993	AF064019	Genbank	3347856
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Sequence 11995	AF064245	Genbank	3930528
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Sequence 12000	AF070646	Genbank	3283920
Sequence 12002	AF070649	Genbank	3283923
Sequence 12003	AF070652	Genbank	4454679
Sequence 12004	AF070655	Genbank	4454685
Sequence 12004 Sequence 12005	AF073792	Genbank	3300101
	AF075053	Genbank	3377594
Sequence 12006 Sequence 12007	AF075242	Genbank	4377992
Sequence 1000	AF077042	Genbank	4689131
Sequence 12008	AF077043	Genbank	4689133
Sequence 12009	AF077046	Genbank	4689139
Sequence 12010	AF077052	Genbank	4689151
Sequence 12011	AF077202	Genbank	4679017
Sequence 12012	AF078855	Genbank	5531824
Sequence 12013	AF081282	Genbank	4336324

Sequence 12014	AF083441	Genbank	5813822
Sequence 12015	AF085361	Genbank	5114056
Sequence 12016	AF086168	Genbank	3483513
Sequence 12017	AF086169	Genbank	3483514
Sequence 12018	AF086210	Genbank	3483555
Sequence 12019	AF086236	Genbank	3483581
Sequence 12020	AF086329	Genbank	3483674
Sequence 12021	AF086465	Genbank	3483810
Sequence 12022	AF086557	Genbank	3483902
Sequence 12023	AF091075	Genbank	3859987
Sequence 12024	AF091083	Genbank	3860003
Sequence 12025	AF091084	Genbank	3860005
Sequence 12026	AF092124	Genbank	3659900
Sequence 12027	AF092135	Genbank	5138919
Sequence 12028	AF098162	Genbank	3929582
Sequence 12029	AF098865	Genbank	4204674
Sequence 12030	AF100748	Genbank	5410281
Sequence 12031	AF100928	Genbank	4323586
Sequence 12032	AF103801	Genbank	6048969
Sequence 12033	AF105278	Genbank .	4324471
Sequence 12034	AF107893	Genbank	4008128
Sequence 12035	AF109872	Genbank	4028621
Sequence 12036	AF110304	Genbank	7106722
Sequence 12037	AF110460	Genbank	4324698
Sequence 12038	AF110774	Genbank	6523794
Sequence 12039	AF111847	Genbank	7211441
Sequence 12040	AF112215	Genbank	6563217
Sequence 12041	AF112222	Genbank	6563229
Sequence 12042	AF113016	Genbank	6642755
Sequence 12043	AF113680	Genbank	6855605
Sequence 12044	AF113700	Genbank	6855634
Sequence 12045	AF117235	Genbank	6563243
Sequence 12046	AF117829	Genbank	4151947
Sequence 12047	AF118091 :	Genbank	6650827
Sequence 12048	AF119663	Genbank	6563251
Sequence 12049	AF121219	Genbank	4903275
Sequence 12050	AF122004	Genbank	5059061
Sequence 12051	AF128536	Genbank	5305705
Sequence 12052	AF129927	Genbank	4558506
Sequence 12053	AF130249	Genbank	4455134
Sequence 12054	AF131738	Genbank	4406548
Sequence 12055	AF131838	Genbank	4406677
Sequence 12056	AF131856	Genbank	4406702
Sequence 12057	AF132965	Genbank	4680700
Sequence 12058	AF132973	Genbank	4680716
Sequence 12059	AF139065	Genbank	6635330
Sequence 12060	AF144700	Genbank	5107075
Sequence 12061	AF144755	Genbank	5006628
Sequence 12062	AF146191	Genbank	5678818
Sequence 12063	AF147330	Genbank	4761681
Sequence 12064	AF150089	Genbank	5107165
Sequence 12065	AF151020	Genbank	7106761
Sequence 12066	AF151066	Genbank	7106853
Sequence 12087	AF151109	Genbank	6166337
Sequence 12068	AF151807	Genbank	4929566
Sequence 12069	AF151832	Genbank	4929616
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Sequence 12070	AF151841	Genbank	4929634
Sequence 12071	AF151861	Genbank	4929674
Sequence 12072	AF151875	Genbank	4929702
Sequence 12073	AF151879	Genbank	4929710
Sequence 12074	AF151884	Genbank	4929720
Sequence 12075	AF151886	Genbank	4929724
Sequence 12076	AF151889	Genbank	4929730
Sequence 12077	AF151893	Genbank	4929738
Sequence 12078	AF151905	Genbank	4929762
Sequence 12079	AF151906	Genbank	4929764
Sequence 12080	AF152462	Genbank	5565976
Sequence 12081	AF155568	Genbank	5031511
Sequence 12082	AF156965	Genbank	5731112
Sequence 12083	AF161389	Genbank	6841191
Sequence 12084	AF161448	Genbank	6841309
Sequence 12085	AF161466	Genbank	6841455
Sequence 12086	AF161500	Genbank	6841523
Sequence 12087	AF161503	Genbank	6841529
Sequence 12088	AF161512	Genbank	6841547
Sequence 12089	AF161556	Genbank	6841379
Sequence 12090	AF165926	Genbank	5817858
Sequence 12091	AF168787	Genbank	7239175
Sequence 12092	AF168956	Genbank	5702387
Sequence 12093	AF174496	Genbank	7108914
Sequence 12094	AF176574	Genbank	5762481
Sequence 12095	AF179274	Genbank	5911402
Sequence 12096	AF187320	Genbank	6164847
Sequence 12097	AF187554	Genbank	6653225
Sequence 12098	AF188178	Genbank	6164864
Sequence 12099	AF188745	Genbank	6425043
Sequence 12100	AF191018	Genbank	6457339
Sequence 12101	AF191298	Genbank	7656642
Sequence 12102	AF196969	Genbank	6289073
Sequence 12103	AF201077	Genbank	6456748
Sequence 12104	AF203815	Genbank	6979641
Sequence 12105	AF205588	Genbank	6531 <b>675</b>
Sequence 12106	AF205600	Genbank	6581094
Sequence 12107	AF207550	Genbank	6470333
Sequence 12108	AF217403	Genbank	6691160
Sequence 12109 Sequence 12110	AF224669	Genbank	7012905
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Sequence 12113	AJ010341 AJ011803	Genbank	4490794
Sequence 12114	AJ012499	Genbank	4753760
Sequence 12115	AJ131693	Genbank	5441359
Sequence 12116	AJ 224442	Genbank	4584422
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Sequence 12118	AJ243009 AJ249731	Genbank	6688156
Sequence 12119	AJ250042	Genbank	5921469
Sequence 12120	AJ250042 AJ250915	Genbank	6013005
Sequence 12121	AK000049	Genbank Genbank	6996445
Sequence 12122	AK000049 AK000052		7019880
Sequence 12123	AK000052 AK000068	Genbank Genbank	7019885
Sequence 12124	AK000083	Genbank Genbank	7019914
Sequence 12125	AK000083 AK000171	Genbank	7019940
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Sequence 12126	AK000221	Genbank	7020163
Sequence 12127	AK000303	Genbank	7020295
Sequence 12128	AK000319	Genbank	7020324
Sequence 12129	AK000395	Genbank	7020455
Sequence 12130	AK000400	Genbank	7020463
Sequence 12131	AK000434	Genbank	7020520
Sequence 12132	AK000462	Genbank	7020566
Sequence 12133	AK000470	Genbank	7020580
Sequence 12134	AK000526	Genbank	7020680
Sequence 12135	AK000560	Genbank	7020738
Sequence 12136	AK000615	Genbank	7020827
Sequence 12137	AK000620	Genbank	7020834
Sequence 12138	AK000624	Genbank	7020839
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Sequence 12140	AK000667	Genbank	7020906
Sequence 12141	AK000669	Genbank	7020909
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Sequence 12143	AK000931	Genbank	7021906
Sequence 12144	AK001067	Genbank	7022104
Sequence 12145	AK001088	Genbank	7022136
Sequence 12146	AK001163	Genbank	7022244
Sequence 12147	AK001192	Genbank	7022292
Sequence 12148	AK001221	Genbank	7022340
Sequence 12149	AK001321	Genbank	7022503
Sequence 12150	AK001338	Genbank	7022533
Sequence 12151	AK001354	Genbank	7022561
Sequence 12152	AK001408	Genbank	7022645
Sequence 12153	AK001441	Genbank	7022699
Sequence 12154	AK001453	Genbank	7022720
Sequence 12155	AK001489	Genbank	7022777 7022808
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Sequence 12159	AK001614	Genbank	7023104
Sequence 12160	AK001689	Genbank	7023153
Sequence 12161	AK001718	Genbank	7023235
Sequence 12162	AK001764 AK001780	Genbank	7023265
Sequence 12163	AK001780 AK001902	Genbank	7023455
Sequence 12164	AK002031	Genbank	7023667
Sequence 12165	AK002001	Genbank	7023777
Sequence 12166	AL002100 AL008721	Genbank	3171883
Sequence 12167 Sequence 12168	AL008723	Genbank	5101762
Sequence 12169	AL009179	Genbank	3217024
Sequence 12170	AL020993	Genbank	3980107
Sequence 12171	AL020997	Genbank	2804155
Sequence 12172	AL021368	Genbank	3080468
Sequence 12173	AL021397	Genbank	3319943
Sequence 12174	AL021707	Genbank	4582132
Sequence 12175	AL021917	Genbank	9367203
Sequence 12176	AL022153	Genbank	3319674
Sequence 12177	AL022238	Genbank	4176442
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Sequence 12179	AL022313	Genbank	4200326
Sequence 12180	AL022326	Genbank	3550039
Sequence 12181	AL022718	Genbank	3763969

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Sequence 12182		Genbank	5679748
Sequence 12183	AL023581	Genbank	3445283
Sequence 12184		Genbank	3805923
Sequence 12185	AL023803	Genbank	9408726
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Sequence 12187	AL024474	Genbank	
Sequence 12188	AL031005	Genbank	3395511
Sequence 12189		Genbank	3287156
Sequence 12190	AL031274	Genbank	3738133
Sequence 12191	AL031295	Genbank	3421083
Sequence 12192	AL031297	Genbank	4376011
Sequence 12193	AL031427	Genbank	4902714
Sequence 12194	AL031432	Genbank	4835258
Sequence 12195	AL031665	Genbank	4375969
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Sequence 12197	AL031718	Genbank	9408729
Sequence 12198	AL031774	_	5805110
Sequence 12199	AL033377	Genbank	3881948
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Sequence 12202	AL033517	Genbank	4140315
Sequence 12203	AL033531	Genbank	4191258
Sequence 12204	AL033531 AL034548	Genbank	6807582
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Sequence 12209		Genbank	6136940
Sequence 12210	AL035413	Genbank	6010110
Sequence 12211	AL035454	Genbank	9368364
Sequence 12212	AL035458	Genbank	6624641
Sequence 12213	AL035461	Genbank	5123778
Sequence 12214	AL035541	Genbank	9581764
Sequence 12215	AL035610	Genbank	6249454
Sequence 12216	AL035699	Genbank	4826515
Sequence 12217	AL049280	Genbank	4500037
Sequence 12218	AL049294	Genbank	4500055
Sequence 12219	AL049447	Genbank	4500230
Sequence 12220	AL049461	Genbank	4500250
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Sequence 12224	AL049694	Genbank	5650653
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Sequence 12226	AL049766	Genbank	5763746
Sequence 12227	AL049779	Genbank	8176894
Sequence 12228	AL049829	Genbank	8217859
Sequence 12229	AL049844	Genbank	5924019
Sequence 12230	AL049849	Genbank	4826529
Sequence 12231	AL049869	Genbank	8217861
Sequence 12232	AL049870	Genbank	7248268
Sequence 12233	AL049911	Genbank	6114591
Sequence 12234	AL049944	Genbank	4884189
Sequence 12235	AL049962	Genbank	4884212
Secuence 12235	AL049964	Genbank	4884214
Sequence 12236 Sequence 12237	AL049968	Genbank	4884217
	AL050016	Genbank	4884262

Sequence 12238	AL050089	Genbank	4884107
Sequence 12239	AL050275	Genbank	4886500
Sequence 12240	AL050277	Genbank	4886454
Sequence 12241	AL050343	Genbank	6137008
Sequence 12242	AL078463	Genbank	5777575
Sequence 12243	AL078621	Genbank	6013067
Sequence 12244	AL079333	Genbank	5738434
Sequence 12245	AL080080	Genbank	5262491
Sequence 12246	AL080102	Genbank	5262526
Sequence 12247	AL080193	Genbank	5262674
Sequence 12248	AL080276	Genbank ·	• • • • • •
Sequence 12249	AL109672	Genbank	5689836
Sequence 12250	AL109759	Genbank	8176900
Sequence 12251	AL109798	Genbank	6453386
Sequence 12252	AL109865	Genbank	6911646
Sequence 12253	AL110183	Genbank	5817095
Sequence 12254	AL110185	Genbank	5817098
Sequence 12255	AL110193	Genbank	5817109
Sequence 12256	AL110194	Genbank	5817111
Sequence 12257	AL110202	Genbank	5817121
Sequence 12258	AL110271	Genbank	5817083
Sequence 12259	AL117350	Genbank	7159817
Sequence 12260	AL117450	Genbank	5911900
Sequence 12261	AL117461	Genbank	5911922
Sequence 12262	AL117482	Genbank	5911960
Sequence 12263	AL117648	Genbank	5912240
Sequence 12264	AL117662	Genbank	5912257 6434634
Sequence 12265	AL121603	Genbank	
Sequence 12266	AL121652	Genbank	7159615
Sequence 12267	AL121716	Genbank	7159797 8919824
Sequence 12268	AL121790	Genbank	6635879
Sequence 12269	AL121823	Genbank	8176919
Sequence 12270	AL121852	Genbank	7161783
Sequence 12271	AL121963	Genbank	6093222
Sequence 12272	AL122036	Genbank Genbank	6102860
Sequence 12273	AL122066	<del>-</del>	6453416
Sequence 12274	AL133010	Genbank Genbank	6453483
Sequence 12275	AL133060	Genbank	6453507
Sequence 12276	AL133069	Genbank	7228177
Sequence 12277	AL133215	Genbank	6983473
Sequence 12278	AL133216	Genbank	7076389
Sequence 12279	AL133387 AL133453	Genbank	7009594
Sequence 12280	AL133593	Genbank	6599181
Sequence 12281	AL133662	Genbank	6599256
Sequence 12282		Genbank	6682291
Sequence 12283	AL135744	Genbank	6635874
Sequence 12284	AL135959 AL136296	Genbank	8217913
Sequence 12285		Genbank	7018294
Sequence 12286	AL136504	Genbank	6807666
Sequence 12287	AL136831 AL137269	Genbank	6807703
Sequence 12288		Genbank	6808024
Sequence 12289	AL137450	Genbank	6808031
Sequence 12290	AL137454	Genbank	6808096
Sequence 12291	AL137481 AL137542		6808221
Sequence 12292		Genbank	6807841
Sequence 12293	AL137673	Golibalik	550.511

Sequence 12294	AL137714	Genbank	6808128
Sequence 12295	AL137721	Genbank	6808159
Sequence 12296	AL137723	Genbank	6808203
Sequence 12297	AL137818	Genbank	7009598
Sequence 12298	AL157442	Genbank	7018558
Sequence 12299	AL157468	Genbank	7018484
Sequence 12300	AL157493	Genbank	7018539
Sequence 12301	AL158493	Genbank	7161641
Sequence 12302	AL158547	Genbank	7161742
Sequence 12303	AP000008	Genbank	3132318
Sequence 12304	AP000009	Genbank	4666255
Sequence 12305	AP000037	Genbank	3132347
Sequence 12306	AP000038	Genbank	
Sequence 12307	AP000346	Genbank	3132348
Sequence 12308	AP000347	Genbank	5103009
Sequence 12309	AP000473	Genbank	5103010
Sequence 12310	AP000497	Genbank	6942329
Sequence 12311	AP000525	Genbank	5926684
Sequence 12312	AP000547	Genbank	5931503
Sequence 12313	AP000692	Genbank	5931525
Sequence 12314	AP000962	Genbank	6693635
Sequence 12315	AP001063		6942330
Sequence 12316	AP001115	Genbank	6693613
Sequence 12317	AP001136	Genbank	6899854
Sequence 12318	AP001137	Genbank	6970360
Sequence 12319	AP001138	Genbank	6970361
Sequence 12320	AP001172	Genbank	7262569
Sequence 12321	AP001343	Genbank	6983884
Sequence 12322	D00596	Genbank	7209831
Sequence 12323	D13119	Genbank	220135
Sequence 12324	D13119	Genbank	285909
Sequence 12325	D13634	Genbank	286026
Sequence 12326	D13654 D14658	Genbank	285992
Sequence 12327	D14665	Genbank	285940
Sequence 12328	D14694	Genbank	6630617
Sequence 12329	D14696	Genbank	603801
Sequence 12330	D14096 D14710	Genbank	285962
Sequence 12331	D16111	Genbank	559324
Sequence 12332	D16947	Genbank	435637
Sequence 12333	D17652	Genbank	598932
Sequence 12334	D23662	Genbank	409069
Sequence 12335	D25274	Genbank	432362
Sequence 12336	D26600	Genbank	464185
Sequence 12337	D29805	Genbank	565650
Sequence 12338	D31885	Genbank	474986
Sequence 12339	D31665 D38112	Genbank	505097
Sequence 12340	D38550	Genbank	644480
Sequence 12341	D36350 D42052	Genbank	559708
Sequence 12342	D42052	Genbank	1136740
Sequence 12343	D42063	Genbank	924266
Sequence 12343	D45131	Genbank	1304103
Sequence 12345	D50916	Genbank	1469174
Sequence 12345	D50929	Genbank	1469200
	D55673	Genbank	870746
Sequence 12347	D63486	Genbank	1469885
Sequence 12348	D78130	Genbank	2443315
Sequence 12349	D79205	Genbank	1754620

10050	D79991	Genbank	1136397
Sequence 12350		Genbank	1136415
Sequence 12351	D80000 D83077	Genbank	1304131
Sequence 12352	D83647	Genbank	1209009
Sequence 12353	D83767	Genbank	1913784
Sequence 12354	D84273	Genbank	3894217
Sequence 12355	D84344	Genbank	2340829
Sequence 12356	D86971	Genbank	1504015
Sequence 12357	D86982	Genbank	1504037
Sequence 12358	D87450	Genbank	1665788
Sequence 12359	D87666	Genbank	1620016
Sequence 12360	D87684	Genbank	1663703
Sequence 12361	D87742	Genbank	1665824
Sequence 12362	E01932	Genbank	2170180
Sequence 12363	E02339	Genbank	2170574
Sequence 12364	E02628	Genbank	2170856
Sequence 12365	E03413	Genbank	2171629
Sequence 12366	E08204	Genbank	2176325
Sequence 12367	E12274	Genbank	3251108
Sequence 12368	E13298	Genbank	3252103
Sequence 12369	E13330	Genbank	3252135
Sequence 12370	E15239	Genbank	5709922
Sequence 12371 Sequence 12372	J00139	Genbank	182715
Sequence 12373	J01415	Genbank	1944628
Sequence 12374	J02960	Genbank	178203
Sequence 12375	J03537	Genbank	337513
Sequence 12376	J03592	Genbank	339722
Sequence 12377	J03746	Genbank	183655
Sequence 12378	J03779	Genbank	179833
Sequence 12379	J04205	Genbank	178686
Sequence 12380	K00558	Genbank	340020
Sequence 12381	L01057	Genbank	292462
Sequence 12382	L05 <b>09</b> 1	Genbank	388030
Sequence 12383	L05092	Genbank	388031
Sequence 12384	L05628	Genbank	1835658
Sequence 12385	L08125	Genbank	306500
Sequence 12386	L11910	Genbank	292420
Sequence 12387	L12350	Genbank	307505 306554
Sequence 12388	L13806	Genbank	292165
Sequence 12389	L13850	Genbank	307437
Sequence 12390	L14076	Genbank	440307
Sequence 12391	L19185	Genbank	306467
Sequence 12392	L19597	Genbank	306741
Sequence 12393	L20046	Genbank	507251
Sequence 12394	L20941	Genbank	348706
Sequence 12395	L22569	Genbank	409465
Sequence 12396	L25616	Genbank	454151
Sequence 12397	L28809	Genbank	1220360
Sequence 12398	L31610	Genbank	532031
Sequence 12399	L35681	Genbank	893287
Sequence 12400	L38951	Genbank	704415
Sequence 12401	L38995	Genbank	1235613
Sequence 12402	L39332	Genbank	1237109
Sequence 12403	L39350	Genbank Genbank	762825
Sequence 12404	L41349	Genbank Genbank	951202
Sequence 12405	L43964	Genbank	551252

Sequence 12406	L47647	Genbank	1000861
Sequence 12407	L76416	Genbank	5566605
Sequence 12408	L76568	Genbank	
Sequence 12409	M11058	Genbank	1905925 184243
Sequence 12410	M11167	Genbank	
Sequence 12411	M11353	Genbank	337381
Sequence 12412	M13519	Genbank	184092
Sequence 12413	M16342	_	179461
Sequence 12414	M16541	Genbank	184266
Sequence 12415	M17885	Genbank	1280204
Sequence 12416	M17886	Genbank	190231
Sequence 12417	M18930	Genbank	190233
Sequence 12418	M19961	Genbank	184371
Sequence 12419	M20496	Genbank	180940
Sequence 12420	M21142	Genbank	809235
Sequence 12421	M22348	Genbank	183402
Sequence 12422		Genbank	190815
Sequence 12423	M22760	Genbank	695359
Sequence 12424	M22919	Genbank	189016
Sequence 12425	M24194	Genbank	187701
Sequence 12426	M24486	Genbank	190785
Sequence 12427	M24543	Genbank	341200
Sequence 12428	M25160	Genbank	189059
Sequence 12429	M26481	Genbank	619789
Sequence 12430	M26663	Genbank	618463
	M27878	Genbank	1020144
Sequence 12431	M30448	Genbank	181154
Sequence 12432	M31520	Genbank	337504
Sequence 12433	M37583	Genbank	184059
Sequence 12434	M38188	Genbank	189378
Sequence 12435	M58485	Genbank	180154
Sequence 12436	M59907	Genbank	189383
Sequence 12437	M60457	Genbank	181249
Sequence 12438	M63005	Genbank	179981
Sequence 12439	M65209	Genbank	184444
Sequence 12440	M74089	Genbank	182400
Sequence 12441	M74775	Genbank	187151
Sequence 12442	M75883	Genbank	432974
Sequence 12443	M77142	Genbank	339700
Sequence 12444	M77836	Genbank	189497
Sequence 12445	M81601	Genbank	339442
Sequence 12446	M90054	Genbank	337579
Sequence 12447	M93036	Genbank	182904
Sequence 12448	M96982	Genbank	338262
Sequence 12449	S62140	Genbank	386156
Sequence 12450	S63912	Genbank	399757
Sequence 12451	S82300	Genbank	245388
Sequence 12452	U02031	Genbank	451329
Sequence 12453	U07151	Genbank	460624
Sequence 12454	U09813	Genbank	1008454
Sequence 12455	U09954	Genbank	607792
Sequence 12456	U12128	Genbank	557287
Sequence 12457	U13616	Genbank	608024
Sequence 12458	U14193	Genbank	555899
Sequence 12459	U14972	Genbank	550024
Sequence 12460	U16798	Genbank	806751
Sequence 12461	U17899	Genbank	717053
			111000

Table 3-5

	- · · -	Cambank	914904
Sequence 12462	U19145	Genbank	736410
Sequence 12463	U21914	Genbank	1215670
Sequence 12464	U24223	Genbank	809486
Sequence 12465	U24231	Genbank	887792
Sequence 12466	U25165	Genbank	799380
Sequence 12467	U25182	Genbank	1045393
Sequence 12468	U25372	Genbank	951331
Sequence 12469	U31657	Genbank	1710107
Sequence 12470	U33886	Genbank	1236349
Sequence 12471	U34683	Genbank	1477465
Sequence 12472	U35246	Genbank	1185451
Sequence 12473	U37143	Genbank	1574941
Sequence 12474	U37230	Genbank	1574947
Sequence 12475	U38784	Genbank	1173904
Sequence 12476	U41371	Genbank	8331754
Sequence 12477	U52111	Genbank	1924973
Sequence 12478	U52427	Genbank	2351381
Sequence 12479	U54562	Genbank	4097314
Sequence 12480	U54993	Genbank	4097314
Sequence 12481	U55184	Genbank	2145094
Sequence 12482	U55853	Genbank	
Sequence 12483	U57847	Genbank	1373420 1438934
Sequence 12484	U61167	Genbank	
Sequence 12485	U65896	Genbank	1763689
Sequence 12486	U66469	Genbank	1724072
Sequence 12487	U66871	Genbank	1519518
Sequence 12488	U68105	Genbank	1562509
Sequence 12489	U68758	Genbank	4097815
Sequence 12490	U69570	Genbank	1825473
Sequence 12491	U70063	Genbank	1743866 2062372
Sequence 12492	U73379	Genbank	1857236
Sequence 12493	U73824	Genbank	1737211
Sequence 12494	U80017	Genbank	2231366
Sequence 12495	U80763	Genbank	1845344
Sequence 12496	U81375	Genbank	6960456
Sequence 12497	U82695	Genbank	1913889
Sequence 12498	U90909	Genbank	2344876
Sequence 12499	U91322	Genbank	2088550
Sequence 12500	U91328	Genbank	1946691
Sequence 12501	U94586	Genbank	2804669
Sequence 12502	U95689	Genbank	28251
Sequence 12503	X00351	Genbank Genbank	34312
Sequence 12504	X02152	Genbank	34309
Sequence 12505	X03083	Genbank	30204
Sequence 12506	X06409	Genbank	30146
Sequence 12507	X15822	Genbank	37495
Sequence 12508	X16064	Genbank	36125
Sequence 12509	X52839		37457
Sequence 12510	X54540	Genbank	509290
Sequence 12511	X60221	Genbank	29508
Sequence 12512	X61123	Genbank	29382
Sequence 12513	X64707	Genbank	31302
Sequence 12514	X65923	Genbank	34634
Sequence 12515	X66401	Genbank	415848
Sequence 12516	X66503	Genbank	36149
Sequence 12517	X67247	Genbank	30143

Sequence 12518         X68271         Genbank         602341           Sequence 12519         X73114         Genbank         402618           Sequence 12520         X74979         Genbank         400462           Sequence 12521         X75682         Genbank         429094           Sequence 12522         X75861         Genbank         429094           Sequence 12523         X76488         Genbank         2385366           Sequence 12525         X81900         Genbank         2274973           Sequence 12526         X84908         Genbank         1502344           Sequence 12527         X98356         Genbank         1770518           Sequence 12528         X99585         Genbank         1770518           Sequence 12530         Y15286         Genbank         34266           Sequence 12531         Y17448         Genbank         5002578           Sequence 12532         Z13009         Genbank         5002578           Sequence 12533         Z14244         Genbank         5002578           Sequence 12536         Z58044         Genbank         695580           Sequence 12537         Z68277         Genbank         1028077           Sequence 12540 <td< th=""><th></th><th></th><th></th><th></th></td<>				
Sequence 12520 X74979 Genbank 400462 Sequence 12521 X75861 Genbank 429094 Sequence 12522 X75861 Genbank 456258 Sequence 12523 X76488 Genbank 434305 Sequence 12524 X80199 Genbank 2385366 Sequence 12526 X84908 Genbank 1502344 Sequence 12526 X84908 Genbank 1502344 Sequence 12527 X98356 Genbank 1502344 Sequence 12528 X95855 Genbank 1770518 Sequence 12529 Y00815 Genbank 2584788 Sequence 12530 Y15286 Genbank 5002578 Sequence 12531 Y17448 Genbank 5002578 Sequence 12532 Z13009 Genbank 31072 Sequence 12533 Z13009 Genbank 30150 Sequence 12534 Z48570 Genbank 30150 Sequence 12535 Z56866 Genbank 1028097 Sequence 12536 Z58044 Genbank 1028097 Sequence 12537 Z68277 Genbank 1130689 Sequence 12538 Z6909 Genbank 1204107 Sequence 12539 Z70233 Genbank 1204107 Sequence 12540 Z73417 Genbank 1235542 Sequence 12541 Z82214 Genbank 1235542 Sequence 12542 Z83313 Genbank 1235542 Sequence 12543 Z83843 Genbank 1235542 Sequence 12544 Z83843 Genbank 1236542 Sequence 12545 Z83843 Genbank 1236542 Sequence 12546 Z83843 Genbank 1304072 Sequence 12546 Z83843 Genbank 1304072 Sequence 12546 Z83843 Genbank 364072 Sequence 12547 Z88812 Genbank 364072 Sequence 12548 Z83831 Genbank 364072 Sequence 12549 Z92544 Genbank 3319673 Sequence 12548 Z85999 Genbank 2326514 Sequence 12550 Z95126 Genbank 3319673 Sequence 12554 Z83851 Genbank 364072 Sequence 12554 Z83851 Genbank 364072 Sequence 12554 Z83851 Genbank 364072 Sequence 12554 S8999 Genbank 3036783 Sequence 12554 S8999 Genbank 364072 Sequence 12556 Septing SET 1445523 Sequence 12557 Septing SET 1445523 Sequence 12558 Septing SET 1445523 Sequence 12556 Septing SET 1445523 Sequence 12557 Septing SET 1445527 Sequence 12558 Septing SET 1445527 Sequence 12559 Septing SET 1445523 Sequence 12550 Septing SET 1445523 Sequence 12556 Septing SET 1485332 Sequence 12568 Septing SET 1485332 Sequence 12569 Sep		X68271	Genhank	602241
Sequence 12521 X74582 Genbank 429094 Sequence 12521 X75881 Genbank 456258 Sequence 12522 X75861 Genbank 456258 Sequence 12524 X80199 Genbank 2385366 Sequence 12525 X81900 Genbank 1502344 Sequence 12527 X98356 Genbank 1502344 Sequence 12528 X99585 Genbank 1770518 Sequence 12529 Y00815 Genbank 2584788 Sequence 12530 Y15286 Genbank 5002578 Sequence 12531 Y17448 Genbank 5002578 Sequence 12531 Y17448 Genbank 30150 Sequence 12532 Z13009 Genbank 30150 Sequence 12533 Z14244 Genbank 30150 Sequence 12535 Z56866 Genbank 1028097 Sequence 12536 Genbank 30150 Sequence 12537 Z68277 Genbank 1028097 Sequence 12538 Z68044 Genbank 1028097 Sequence 12538 Z68044 Genbank 1028097 Sequence 12538 Z68079 Genbank 1204107 Sequence 12538 Z69709 Genbank 1204107 Sequence 12540 Z73417 Genbank 1224592 Sequence 12541 Z82214 Genbank 12235542 Sequence 12542 Z83313 Genbank 1322397 Sequence 12543 Z83843 Genbank 1322397 Sequence 12544 Z88314 Genbank 1322397 Sequence 12545 Z83813 Genbank 1322397 Sequence 12546 Z83313 Genbank 1346072 Sequence 12547 Z84812 Genbank 1364072 Sequence 12548 Z83848 Genbank 2578095 Sequence 12548 Z83848 Genbank 2678095 Sequence 12546 Sequence 12547 Z84812 Genbank 3164072 Sequence 12548 Z83848 Genbank 2678095 Sequence 12549 Z83851 Genbank 3164072 Sequence 12546 Z83851 Genbank 3164072 Sequence 12547 Z84812 Genbank 3164072 Sequence 12548 Z8999 Genbank 265614 Sequence 12550 Z95126 Genbank 3036783 Sequence 12551 Z97053 Genbank 265614 Sequence 12552 Z98749 Genbank 242581 Sequence 12553 Z98749 Genbank 265614 Sequence 12554 Sequence 12555 Sequence 12556 A0005278 EST 1445523 Sequence 12556 A0005278 EST 1448541 Sequence 12566 A022472 EST 1448541 Sequence 12566 A022472 EST 1488501 Sequence 12566 A022540 EST 1489608 Sequence 12568 A026405 EST 1489608 Sequence 12569 A010519 EST 1477545 Sequence 12566 A022540 EST 1489509 Sequence 12566 A026405 EST 1489608 Sequence 12567 A006422 EST 1492901 Sequence 12568 A026405 EST 1492901 Sequence 12569 A006422 EST 1492901 Sequence 12569 A026424 EST 1492901		X73114	Genbank	
Sequence 12521 X75882 Genbank 429094 Sequence 12523 X76488 Genbank 456288 Sequence 12524 X80199 Genbank 2385366 Sequence 12525 X81900 Genbank 2274973 Sequence 12526 X84908 Genbank 1502344 Sequence 12527 X98356 Genbank 1502344 Sequence 12529 Y00815 Genbank 34266 Sequence 12529 Y00815 Genbank 2584788 Sequence 12529 Y00815 Genbank 34266 Sequence 12530 Y15286 Genbank 5002578 Sequence 12531 Y17448 Genbank 3002578 Sequence 12532 Z13009 Genbank 30150 Sequence 12532 Z13009 Genbank 30150 Sequence 12534 Z48570 Genbank 695580 Sequence 12535 Z66866 Genbank 1028097 Sequence 12536 Z58044 Genbank 1028097 Sequence 12537 Z68277 Genbank 1202575 Sequence 12538 Z69709 Genbank 1204107 Sequence 12540 Z73417 Genbank 1322397 Sequence 12541 Z82214 Genbank 1322397 Sequence 12542 Z83313 Genbank 1235542 Sequence 12542 Z83313 Genbank 1322397 Sequence 12542 Z83313 Genbank 1322397 Sequence 12542 Z83313 Genbank 1322397 Sequence 12545 Z83848 Genbank 1322397 Sequence 12545 Z83848 Genbank 1322397 Sequence 12545 Z83848 Genbank 134062 Sequence 12545 Z83848 Genbank 134062 Sequence 12546 Sequence 12546 Genbank 14564 Sequence 12547 Z84812 Genbank 3319673 Sequence 12548 Z83848 Genbank 2578095 Sequence 12549 Z83848 Genbank 3164072 Sequence 12545 Z83851 Genbank 3319673 Sequence 12550 Z95126 Genbank 3319673 Sequence 12551 Z9599 Genbank 2326514 Sequence 12552 Z98749 Genbank 3036783 Sequence 12553 A000222 EST 1445523 Sequence 12554 S9951 Genbank 3036783 Sequence 12555 A001709 EST 147545 Sequence 12556 A001519 EST 1445523 Sequence 12568 A0264695 EST 1488571 Sequence 12568 A0264695 EST 1488571 Sequence 12568 A026400 EST 1492901 Sequence 12569 A016422 EST 1492901 Sequence 12569 A026422 EST 1492913 Sequence 12569 Sequence 12561 EST 1492901 Sequence 12569 A026422 EST 1492901 Sequence 12569 A026422 EST 1492901 Sequence 12569 Sequence 12567 A026422 EST 1492901	Sequence 12520	X74979	Genhank	
Sequence 12522 X75861 Genbank 456258 Sequence 12524 X80199 Genbank 2385366 Sequence 12525 X81900 Genbank 2274973 Sequence 12526 X84908 Genbank 1502344 Sequence 12527 X98356 Genbank 1502344 Sequence 12529 Y08315 Genbank 34266 Sequence 12529 Y08315 Genbank 34266 Sequence 12529 Y08315 Genbank 2584768 Sequence 12530 Y15286 Genbank 5002578 Sequence 12531 Y17448 Genbank 5002578 Sequence 12531 Y17448 Genbank 30150 Sequence 12532 Z13009 Genbank 31072 Sequence 12533 Z14244 Genbank 30150 Sequence 12534 Z48570 Genbank 695580 Sequence 12535 Z56866 Genbank 1028097 Sequence 12536 Z58044 Genbank 1028097 Sequence 12537 Z68277 Genbank 1130689 Sequence 12538 Z69709 Genbank 1204107 Sequence 12539 Z70233 Genbank 1204107 Sequence 12540 Z73417 Genbank 3125542 Sequence 12541 Z82214 Genbank 5911819 Sequence 12542 Z83313 Genbank 122397 Sequence 12542 Z83313 Genbank 122397 Sequence 12544 Z83843 Genbank 5911819 Sequence 12545 Z83851 Genbank 3164072 Sequence 12546 Z83851 Genbank 3164072 Sequence 12548 Sequence 12549 Sequence 12549 Sequence 12549 Sequence 12548 Sequence 12549 Sequence 12550 Sequence 12551 Z97053 Genbank 319673 Sequence 12554 S83851 Genbank 3164072 Sequence 12548 Sepses Genbank 319673 Sequence 12548 Sepses Genbank 319673 Sequence 12550 Z98749 Genbank 3036783 Sequence 12551 Z97053 Genbank 3036783 Sequence 12552 Z98749 Genbank 3036783 Sequence 12554 S8950 Genbank 3036783 Sequence 12555 A0001709 EST 1445523 Sequence 12556 A002222 EST 1448541 Sequence 12558 A002222 EST 1448541 Sequence 12568 A002224 SST 1486571 Sequence 12568 A002247 EST 144550 Sequence 12568 A002427 EST 1486571 Sequence 12568 A026495 EST 1489091 Sequence 12568 A026495 EST 1489091 Sequence 12568 A026495 EST 1499901 Sequence 12568 A026492 EST 1492901 Sequence 12569 Sequence 12569 SST Sequence 12569 SST Sequence 12569 SST SST SST SST SST SST SST SST SST SS	Sequence 12521			
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Sequence 12759	AA432282	EST	2111127
Sequence 12760	AA434140	EST	2114670
Sequence 12761	AA434140 AA434447	EST	2139054
Sequence 12762	AA435671	EST	2139361
Sequence 12763	AA442377	EST	2140585
Sequence 12764	AA442786	EST	2154255
Sequence 12765		EST	2155461
Sequence 12766	AA442878 AA447199	EST	2155553
Sequence 12767		EST	2159864
Sequence 12767 Sequence 12768	AA448292	EST	2161962
Sequence 12769	AA448409	EST	2162079
Sequence 12769 Sequence 12770	AA453183	EST	2166852
Sequence 12771	AA453411	EST	2167080
Sequence 12771	AA453747	EST	2167416
Sequence 12773	AA454187 AA456524	EST	2167856
Sequence 12774		EST	2179100
Sequence 12775	AA458612	EST	2183519
Sequence 12776	AA459528	EST	2184435
Sequence 12777	AA460187 AA460266	EST	2185003
Sequence 12778	AA463256	EST	2185082
Sequence 12779	AA463899	EST	2188140
Sequence 12780	AA465606	EST	2188783
Sequence 12781	AA467764	EST	2191773
Sequence 12782	AA468230	EST	2194298
Sequence 12783	AA468757	EST	2194764
Sequence 12784	AA470337	EST	2195291
Sequence 12785	AA477608	EST	2197646
Sequence 12786	AA4778103	EST	2206242
Sequence 12787	AA478663	EST	2206737
Sequence 12788	AA478669	EST	2207297
Sequence 12789	AA480081	EST	2207303
Sequence 12790		EST	2208232
Sequence 12791	AA483251	EST	2212064
	AA483698	EST	2212511
Sequence 12792	AA486416	EST	2216580
Sequence 12793	AA491100	EST	2220273
Sequence 12794	AA491505	EST	2218288
Sequence 12795	AA493382	EST	2223223
Sequence 12796	AA493522	EST	2223363
Sequence 12797	AA494118	EST	2223959

<u>Table 3-5</u>

Sequence 12798	AA494433	EST	2224220
Sequence 12799	AA496993	EST	2230314
Sequence 12800	AA501429	EST	2236396
Sequence 12801	AA503827	EST	2238794
Sequence 12802	AA505115	EST	2241275
Sequence 12803	AA507605	EST	2244044
Sequence 12804	AA507791	EST	2244230
•	AA51731	EST	2251779
Sequence 12805			2251859
Sequence 12806	AA513447	EST	2253883
Sequence 12807	AA514215	EST	
Sequence 12808	AA514331	EST	2253839
Sequence 12809	AA514555	EST	2254155
Sequence 12810	AA514835	EST	2254435
Sequence 12811	AA514972	EST	2254572
Sequence 12812	AA515971	EST	2255571
Sequence 12813	AA521142	EST	2261685
Sequence 12814	AA522471	EST	2263183
Sequence 12815	AA523897	EST	2264825
Sequence 12816	AA524190	EST	2265118
Sequence 12817	AA525960	EST	2268029
Sequence 12818	AA526064	EST	2268133
Sequence 12819	AA526186	EST	2268255
Sequence 12820	AA526891	EST	2268960
Sequence 12821	AA526943	EST	2269012
Sequence 12822	AA527040	EST	2269109
Sequence 12823	AA528775	EST	2270844
Sequence 12824	AA531540	EST	2274246
Sequence 12825	AA533286	EST	2277382
Sequence 12826	AA533355	EST	2277451
Sequence 12827	AA534137	EST	2278153
Sequence 12828	AA535571	EST	2279824
Sequence 12829	AA541666	EST	2288100
Sequence 12029	AA541813	EST	2288247
Sequence 12830	AA548722	EST	2319004
Sequence 12831		EST	2321985
Sequence 12832	AA551733	EST	2322421
Sequence 12833	AA552169		2325265
Sequence 12834	AA554726	EST	2325733
Sequence 12835	AA555194	EST	2327668
Sequence 12836	AA557191	EST	
Sequence 12837	AA563905	EST	2335544
Sequence 12838	AA564118	EST	2335757
Sequence 12839	AA565885	EST	2337524
Sequence 12840	AA568360	EST	2341414
Sequence 12841	AA568578	EST	2341632
Sequence 12842	AA570171	EST	2344151
Sequence 12843	AA577170	EST	2354644
Sequence 12844	AA577927	EST	2356111
Sequence 12845		EST	2355203
Sequence 12846		EST	2355735
Sequence 12847	AA582370	EST	2359730
Sequence 12848	AA583169	EST	2367778
Sequence 12849		EST	2397735
Sequence 12850		EST	2398139
Sequence 12851	AA588161	EST	2402336
Sequence 12852		EST	2401716
Sequence 12853		EST	2410689
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Sequence 12854	AA599217	EST	0400040
Sequence 12855	AA602418	EST	2432842
Sequence 12856	AA604553	EST	2435914
Sequence 12857	AA608603	EST	2445417
Sequence 12858		EST	2457031
Sequence 12859	AA608990	EST	2457345
Sequence 12860	AA610801		2457418
Sequence 12861	AA613587	EST	2459229
Sequence 12862	AA613637	EST	2464625
Sequence 12863	AA614729	EST	2464675
Sequence 12864	AA626535	EST	2466925
Sequence 12865	AA627229	EST	2538922
Sequence 12866	AA627804	EST	2540273
Sequence 12867	AA627996	EST	2539899
Sequence 12868	AA631218	EST	2539995
Sequence 12869	AA631224	EST	2553829
Sequence 12870	AA631404	EST	2553835
Sequence 12871	AA631684	EST	2554015
Sequence 12872	AA632005	EST	2554295
Sequence 12873	AA632225	EST	2555419
Sequence 12874	AA632328	EST	2555639
Sequence 12875	AA633317	EST	2555742
Sequence 12876	AA633888	EST	2556731
Sequence 12877	AA634120	EST	2557102
Sequence 12878	AA634360	EST	2557334
Sequence 12879	AA635630	EST	2557574
Sequence 12880	AA636029	EST	2559472
Sequence 12881	AA639730	EST	2559871
Sequence 12882	AA640721	EST	2563509
Sequence 12883	AA640779	EST	2565971
Sequence 12884	AA642858	EST	2566029
Sequence 12885	AA643157	EST	2568076
Sequence 12886	AA643611	EST	2568375
Sequence 12887	AA643763	EST	2568829
Sequence 12888	AA649327	EST	2568981
Sequence 12889	AA653201	EST	2575756
Sequence 12890	AA655033	EST	2589372
Sequence 12891	AA657408	EST	2591187
Sequence 12892	AA659388	EST	2593562
Sequence 12893	AA662764	EST	2595542
Sequence 12894	AA669846	EST	2616755
Sequence 12895	AA669862	EST	2631345
Sequence 12896	AA678121	EST	2631361
Sequence 12897	AA678246	EST EST	2658643
Sequence 12898	AA678311	EST	2658768
Sequence 12899	AA678696		2658833
Sequence 12900	AA682980	EST	2659218
Sequence 12901	AA683334	EST	2668871
Sequence 12902	AA687631	EST	2669225
Sequence 12903	AA688365	EST	2674537
Sequence 12904	AA694202	EST	2675271
Sequence 12905	AA699532	EST	2695140
Sequence 12906	AA699712	EST	2702655
Sequence 12907	AA700166	EST	2702675
Sequence 12908	AA701936	EST	2703129
Sequence 12909	AA702410	EST	2705049
	7 02 7 10	EST	2705523

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Sequence 12910	AA702881	EST	2705994
Sequence 12911	AA704096	EST	2714014
Sequence 12912	AA705746	EST	2715664
Sequence 12913	AA707188	EST	2717106
Sequence 12913	AA70B421	EST	2718339
Sequence 12914	AA714951	EST	2727225
Sequence 12915		EST	2740971
Sequence 12916	AA723194	EST	2752521
Sequence 12917	AA731317		2752 <b>577</b>
Sequence 12918	AA731688	EST	2768114
Sequence 12919	AA736880	EST	
Sequence 12920	AA748046	EST	2788004
Sequence 12921	AA749025	EST	2788983
Sequence 12922	AA764734	EST	2815972
Sequence 12923	AA769434	EST	2820672
Sequence 12924	AA770613	EST	2821851
Sequence 12925	AA772280	EST	2824063
Sequence 12926	AA773040	EST	2824611
Sequence 12927	AA775624	EST	2834958
Sequence 12928	AA775824	EST	2835158
Sequence 12920	AA777499	EST	2836978
Sequence 12929	AA778560	EST	2837891
Sequence 12930		EST	2838107
Sequence 12931	AA778776	EST	2848874
Sequence 12932	AA788754	EST	2875479
Sequence 12933	AA806729		2876454
Sequence 12934	AA806878	EST	2877612
Sequence 12935	AA808206	EST	2878673
Sequence 12936	AA809267	EST	
Sequence 12937	AA809700	EST	2879106
Sequence 12938	AA809964	EST	2879370
Sequence 12939	AA810073	EST	2879479
Sequence 12940	AA811224	EST	2880835
Sequence 12941	AA811879	EST	2881490
Sequence 12942	AA813197	EST	2883182
Sequence 12943	AA814246	EST	2883842
Sequence 12944	AA814643	EST	2884239
Sequence 12945	AA825680	EST	2898992
Sequence 12946	AA828003	EST	2900366
Sequence 12947	AA828330	EST	2901429
Sequence 12948	AA829737	EST	2902836
Sequence 12949	AA831926	EST	2905025
Sequence 12950	AA831948	EST	2905047
Sequence 12951	AA832521	EST	2898667
Sequence 12952	AA836590	EST	2910909
Sequence 12953	AA837128	EST	2912327
Sequence 12954	AA843115	EST	2929633
Sequence 12955	AA843807	EST	2930258
Sequence 12956	AA846016	EST	2932156
Sequence 12957	AA846735	EST	2932875
Sequence 12937	AA854511	EST	2942049
Sequence 12958	AA861279	EST	2953419
Sequence 12959		EST	2955709
Sequence 12960	AA863230	EST	2955856
Sequence 12961	AA863377		2957845
Sequence 12962	AA865569	EST	2968038
Sequence 12963	AA872000	EST	2969295
Sequence 12964	AA873173	EST	2969293 2969611
Sequence 12965	AA873489	EST	2303011

12966	AA876033	EST	2984796
	AA877009	_	2986086
	AA877239		2986316
	AA877769		2986734
	AA883195		2994671
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	· · ·		2994374
			3001182
			3017221
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			3038097
			3038424
			3049981
			3050171
			3057272
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			3089998
			3092853
			3095518
			3109991
			3110585
			3109929
			3118158
	· · · · · · · · ·		3144675
	•		3145479
			3147913
			3148350
		•	3161982
			3172621
			3174559
			3179398
			3179505
			3182007
			3203728
			3215164
			3231245
			3232474
			3241469
			3253790
			3255413
			3279158
			3280033
			3280865
			3281007
			3308126
			3308247
			3308752
			3321531
			3331473
			6358883
			6358939
			6358963
			6359053
			3400787
3021	AIU/4253	EST	3400897
	12966 12967 12968 12969 12970 12971 12972 12973 12974 12975 12976 12977 12980 12981 12982 12983 12984 12985 12986 12987 12988 12999 12999 12999 12999 12999 12999 12999 12999 13000 13001 13002 13003 13004 13005 13006 13007 13008 13007 13008 13009 13011 13008 13009 13011 13012 13013 13014 13015 13016 13017 13018 13019 13019 13019 13019 13019 13019 13019 13017 13018 13019 13019 13020 13020 13020 13020 13020 13020	12967 AA877009 12968 AA877239 12969 AA877769 12970 AA883195 12971 AA884623 12972 AA885297 12973 AA886074 12974 AA890342 12975 AA894745 12976 AA902974 12977 AA903301 12978 AA910691 12979 AA910881 12980 AA917382 12981 AA931246 12982 AA931567 12983 AA93730 12984 AA935696 12985 AA937407 12986 AA946596 12987 AA946596 12989 AA954463 12990 AA969495 12991 AA969966 12989 AA954463 12990 AA969495 12991 AA969966 12992 AA972733 12993 AA973170 12994 AA983457 12995 AA987257 12996 AA988988 12997 AA992853 12998 AA992960 12999 AA995518 13000 AI003314 13001 AI005654 13002 AI016909 13003 AI018676 13004 AI025856 13005 AI032664 13006 AI034460 13007 AI039964 13008 AI041671 13010 AI052135 13011 AI052135 13011 AI052761 13014 AI052761 13015 AI057607 13016 AI061642 13018 AI064691 13019 AI064781 13020 AI074143	12967 AA877009 EST 12968 AA877239 EST 12969 AA877769 EST 12970 AA883195 EST 12971 AA884623 EST 12972 AA885297 EST 12973 AA886074 EST 12974 AA890342 EST 12975 AA994745 EST 12976 AA902974 EST 12977 AA903301 EST 12978 AA910691 EST 12979 AA910881 EST 12980 AA917382 EST 12981 AA931246 EST 12982 AA931567 EST 12983 AA93730 EST 12984 AA935696 EST 12985 AA937407 EST 12986 AA946596 EST 12987 AA946596 EST 12988 AA946676 EST 12989 AA954463 EST 12989 AA954463 EST 12990 AA969495 EST 12991 AA969966 EST 12992 AA973170 EST 12993 AA973170 EST 12994 AA983457 EST 12995 AA987257 EST 12996 AA988988 EST 12997 AA98988 EST 12999 AA992853 EST 12999 AA992853 EST 13000 AI003314 EST 13000 AI003664 EST 13000 AI003740 EST 13000 AI00381676 EST 13000 AI003816 EST 13000 AI005654 EST 13000 AI005654 EST 13000 AI005654 EST 13000 AI003814 EST 13000 AI005654 EST 13000 AI005654 EST 13000 AI005656 EST 13000 AI005656 EST 13000 AI005256 EST 13000 AI005256 EST 13000 AI005256 EST 13001 AI052256 EST 13001 AI052256 EST 13001 AI052256 EST 13013 AI052266 EST 13014 AI052761 EST 13015 AI057607 EST 13016 AI061642 EST 13017 AI061642 EST 13018 AI064691 EST 13019 AI064781 EST 13020 AI074143 EST

**Table 3-5** 

Sequence 13022	A1074350	EST	3400994
Sequence 13023	AI075208	EST	3401799
Sequence 13024	AI076009	EST	3405187
Sequence 13025	AI076793	EST	3405971
Sequence 13026	AI079885	EST	3416136
Sequence 13027	AI080676	EST	3416927
Sequence 13028	Al081536	EST	3418328
Sequence 13029	A1086009	EST	3424432
Sequence 13030	AI086963	EST	3425386
Sequence 13031	AI087104	EST	3425527
Sequence 13032	AI091297	EST	3430356
Sequence 13033	AI091340	EST	3430399
Sequence 13034	AI096751	EST	3445859
Sequence 13035	Al114651	EST	6359996
Sequence 13036	Al123912	EST	3539678
Sequence 13037	Al126019	EST	3594533
Sequence 13038	Al127959	EST	3596473
Sequence 13039	Al133330	EST	6360646
Sequence 13040	Al139392	EST	3645364
Sequence 13041	Al143894	EST	3665703
Sequence 13042	AI144216	EST	3666025
Sequence 13043	AI146737	EST	3674419
Sequence 13044	AI148343	EST	3676025
Sequence 13045	AI159872	EST	3693252
Sequence 13046	AI160110	EST	3693490
Sequence 13047	AI168423	EST	3701593
Sequence 13048	Al183462	EST	3734100
Sequence 13049	Al184770	EST	3735408
Sequence 13050	Al186840	EST	3737478
Sequence 13051	Al187815	EST	3739024
Sequence 13052	Al189478	EST	3740687
Sequence 13053	Al191342	EST	3742551
Sequence 13054	Al198812	EST	3751418
Sequence 13055	Al198882	EST	3751488
Sequence 13056	Al202429	EST	3755035
Sequence 13057	Al202937	EST	3755543
Sequence 13058	Al207682	EST	6361696
Sequence 13059	Al216387	EST	3785428
Sequence 13060	Al221087	EST .	3803290
Sequence 13061	Al240307	EST	3835704
Sequence 13062	Al240817	EST	3836214
Sequence 13063	Al242075	EST	3837472
Sequence 13064	Al244187	EST	3839584
Sequence 13065	Al246248	EST	3841645
Sequence 13066	AI247411	EST	3842808
Sequence 13067			3843922
	AI248525	EST	
Sequence 13068	AI249008	EST	3844405
Sequence 13069	AI251793	EST	3848322
Sequence 13070	AI253046	EST	3849575
Sequence 13071	AI253330	EST	3850451
Sequence 13072	AI253436	EST	3850391
Sequence 13073	AI262640	EST	3870843
Sequence 13074	AI267506	EST	3886673
Sequence 13075	AI267570	EST	3886737
Sequence 13076	AI268542	EST	3887709
Sequence 13077	Al269205	EST	3888372

Sequence 13078	Al275967	EST	3898241
Sequence 13079	Al275998	EST	3898272
Sequence 13080	Al276110	EST	3898384
Sequence 13081	Al278317	EST	3900585
Sequence 13082	Al278766	EST	3917000
Sequence 13083	Al278998	EST	3917232
Sequence 13084	Al280044	EST	3918277
Sequence 13085	Al280747	EST	
Sequence 13086	Al281464		3918980
Sequence 13087	Al282246	EST EST	3919697
Sequence 13088	Al282254	EST	3920479
Sequence 13089	Al285371		3920487
Sequence 13090	Al288305	EST	3923604
Sequence 13091	Al289108	EST	3930985
Sequence 13092	Al290733	EST	3932372
		EST	3933507
Sequence 13093	AI291613	EST	3934387
Sequence 13094	Al291918	EST	3934692
Sequence 13095	AI302808	EST	3962154
Sequence 13096	Al309128	EST	4003999
Sequence 13097	AI335458	EST	4072385
Sequence 13098	Al337195	EST	4074122
Sequence 13099	Al339940	EST	4076867
Sequence 13100	Al341735	EST	4078662
Sequence 13101	Al342244	EST	4079171
Sequence 13102	Al344445	EST	4081651
Sequence 13103	A1345345	EST	4082551
Sequence 13104	AI345677	EST	4082883
Sequence 13105	AI356050	EST	4107671
Sequence 13106	AI361513	EST	4113134
Sequence 13107	Al370743	EST	4149496
Sequence 13108	Al380670	EST	4190523
Sequence 13109 Sequence 13110	Al381627 Al384017	EST	4194408
Sequence 13111	Al400359	EST	4196798
Sequence 13112	Al420311	EST EST	4243446
Sequence 13113	AI421401	EST	4266242
Sequence 13114	Al435207	EST	4267332 4301566
Sequence 13115	Al446753	EST	4297414
Sequence 13116	Al458730	EST	4311309
Sequence 13117	Al458971	EST	4311550
Sequence 13118	Al459103	EST	4311682
Sequence 13119	Al459338	EST	4311917
Sequence 13120	Al468004	EST	4330094
Sequence 13121	Al469505	EST	4331595
Sequence 13122	Al479751	EST	4372919
Sequence 13123	Al480118	EST	4373286
Sequence 13124	Al492144	EST	4393147
Sequence 13125	Al497917	EST	4389899
Sequence 13126	Al499678	EST	4391660
Sequence 13127	Al500313	EST	4392295
Sequence 13128	AI554818	EST	4487181
Sequence 13129	Al559531	EST	4509736
Sequence 13130	AI560688	EST	4509736 4511029
Sequence 13131	AI566007	EST	4511029 4524459
Sequence 13132	Al567612	EST	4524459 4526064
Sequence 13133	AI568146	EST	4526720
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**Table 3-5** 

Sequence 1	13134	AI568614	EST	4531988
Sequence 1	13135	AI572600	EST	4535974
Sequence 1		AI581048	EST	4565424
Sequence 1		AI582231	EST	4568128
Sequence 1		AI58B871	EST	4597919
			EST	4606830
Sequence :		AI597782		
Sequence 1		Al610953	EST	4620120
Sequence '		Al611995	EST	4621162
Sequence 1	13142	Al619754	EST	4628880
Sequence :	13143	AI627855	EST	4664655
Sequence 1	13144	Al631608	EST	4682938
Sequence 1	13145	AI631996	EST	4683326
Sequence 1		Al632109	EST	4683439
Sequence 1		A1632955	EST	4684285
Sequence '		Al636794	EST	4688124
Sequence :		Al653101	EST	4737080
		Al653441	EST	4737420
Sequence :			EST	4739623
Sequence 1		Al655644		
Sequence :		AI659251	EST	4762821
Sequence 1		Al659386	EST	4762956
Sequence 1		Al669708	EST	4834482
Sequence 1	13155	AI678630	EST	4888812
Sequence 1	13156	Al681539	EST	4891721
Sequence 1		AI683912	EST	4894094
Sequence '		Al684760	EST	4896054
Sequence '		Al685829	EST	4897123
Sequence '		Al688152	EST	4899446
Sequence :		Al688730	EST	4900024
Sequence :		Al690281	EST	4901575
Sequence		A1692476	EST	4969816
Sequence 1		Al692763	EST	4970103
Sequence :		Al694509	EST	4971849
Sequence '		AI695805	EST	4983705
Sequence '	13167	AI703457	EST	4991357
Sequence :	13168	AI741323	EST	5109611
Sequence :	13169	AI743098	EST	5111386
Sequence '	13170	AI744238	EST	5112526
Sequence 1		AI744256	EST	5112544
Sequence :		Al744984	EST	5113272
Sequence '		Al749216	EST	5127480
Sequence '		Al752250	EST	5130514
Sequence '		Al753825	EST	5132177
Sequence :		Al760744	EST	5176411
			EST	5178544
Sequence		Al762877		
Sequence '		Al766072	EST	5232581
Sequence 1		Al769461	EST	5235970
Sequence '		Al792057	EST	5339773
Sequence :	13181	Al796688	EST	5362151
Sequence :	13182	AI802907	EST	5368368
Sequence :	13183	Al803019	EST	5368491
Sequence :		AI803566	EST	5369026
Sequence '		Al806628	EST	5393194
Sequence 1		AI811625	EST	5398191
Sequence		AI815404	EST	5430950
		Al815499	EST	5431045
Sequence :				-
Sequence :	13789	Al816161	EST	5431707

**Table 3-5** 

Sequence 13190	Al819909	EST	5438988
Sequence 13191	Al824193	EST	5444864
Sequence 13192	A1826628	EST	5447299
Sequence 13193	AI859220	EST	5512836
Sequence 13194	AI863349	EST	5527456
Sequence 13195	Al866561	EST	5530668
Sequence 13196	AI868330	EST	5541346
Sequence 13197	Al872857	EST	5546906
Sequence 13198	Al889718	EST	5594882
Sequence 13199	Al905192	EST	6495579
Sequence 13200	Al912093	EST	5631948
Sequence 13201	Al921575	EST	5657539
Sequence 13202	Al922278	EST	5658242
Sequence 13203	Al925156	EST	5661120
Sequence 13204	Al928710	EST	5664674
Sequence 13205	Al932970	EST	5671707
Sequence 13206	Al939359	EST	5678229
Sequence 13207	Al951970	EST	5744280
Sequence 13208	Al952576	EST	5744886
Sequence 13209	Al954275	EST	5746585
Sequence 13210	Al962421	EST	5755134
Sequence 13211	Al962746	EST	5755459
Sequence 13212	Al968255	EST	5765073
Sequence 13213	Al970812	EST	5767638
Sequence 13214	AI991063	EST	5837966
Sequence 13215	Al992283	EST	5839188
Sequence 13216	AL036115	EST	5927646
Sequence 13217	AL036349	EST	5405930
Sequence 13218	AL036413	EST	5405969
Sequence 13219	AL037368	EST	5406772
Sequence 13220	AL038505	EST	5928459
Sequence 13221	AL040528	EST	5935315
Sequence 13222	AL040985	EST	5409929
Sequence 13223	AL042384	EST	5421723
Sequence 13224	AL044640	EST	5432855
Sequence 13225	AL048351	EST	5936451
Sequence 13226	AL134733	EST	6602920
Sequence 13227	AL157600	EST	7058001
Sequence 13228	AW004978	EST	5853682
Sequence 13229	AW014893	EST	5863650
Sequence 13230	AW020158	EST	5873698
Sequence 13231	AW027717	EST	5886473
Sequence 13232	AW051707	EST	5913989
Sequence 13233	AW069100	EST	6024098
Sequence 13234	AW069166	EST	6024252
Sequence 13235	AW069188	EST	6024274
Sequence 13236	AW070930	EST	6025928
Sequence 13237	AW083185	EST	6038337
Sequence 13238	AW084353	EST	6039505
Sequence 13239	AW086188	EST	6041262
Sequence 13240	AW088595	EST	6044412
Sequence 13241	AW089333	EST	6046677
Sequence 13242	AW118097	EST	6086681
Sequence 13243	AW136562	EST	6140695
Sequence 13244	AW 130302 AW 148690	EST	6196586
Sequence 13245	AW 151847	EST	6199832
		LUI	0100002

Sequence 13246	AW166819	EST	6398344
Sequence 13247	AW167129	EST	6398654
Sequence 13248	AW168050	EST	6399575
Sequence 13249	AW178252	EST	6444289
Sequence 13250	AW188275	EST	6462711
Sequence 13251	AW188668	EST	6463028
Sequence 13252	AW 100000 AW 190301	EST	6464781
	AW192683	EST	6471382
Sequence 13253			6472756
Sequence 13254	AW194025	EST	6473323
Sequence 13255	AW194497	EST	6474734
Sequence 13256	AW195584	EST	
Sequence 13257	AW196723	EST	6475953
Sequence 13258	AW204679	EST	6504151
Sequence 13259	AW207172	EST	6506668
Sequence 13260	AW243325	EST	6577165
Sequence 13261	AW263682	EST	6640498
Sequence 13262	AW290969	EST	6697605
Sequence 13263	AW292970	EST	6699606
Sequence 13264	AW295194	EST	6701830
Sequence 13265	AW297219	EST	6703855
Sequence 13266	AW301845	EST	6711522
Sequence 13267	AW302764	EST	6712444
Sequence 13268	AW364986	EST	6869636
Sequence 13269	AW365134	EST	6869784
Sequence 13270	AW391931	EST	6896590
Sequence 13271	AW402563	EST	6921261
Sequence 13272	AW405635	EST	6924692
Sequence 13273	AW410936	EST	6936477
Sequence 13274	AW411275	EST	6936816
Sequence 13275	AW444608	EST	6986370
Sequence 13276	AW451259	EST	6992035
Sequence 13277	AW474887	EST	7044993
Sequence 13278	AW578558	EST	7253607
Sequence 13279	F29317	EST	4814943
Sequence 13280	F34866	EST	4820492
Sequence 13281	F36899	EST	4822525
Sequence 13282	H17774	EST	884014
Sequence 13283	H54240	EST	994387
Sequence 13284	H56010	EST	1004654
Sequence 13285	H71857	EST	1043673
Sequence 13286	H78871	EST	1056960
Sequence 13287	H97996	EST	1118881
Sequence 13288	N20525	EST	1125480
	N27572	EST	1142053
Sequence 13289		EST	1190084
Sequence 13290	N48918		
Sequence 13291	N62491	EST	1210320
Sequence 13292	N95724	EST	1268009
Sequence 13293	R02774	EST	752510
Sequence 13294	R06525	EST	757145
Sequence 13295	R14391	EST	767467
Sequence 13296	R55374	EST	824669
Sequence 13297	T08334	EST	389362
Sequence 13298	T24823	EST	534448
Sequence 13299	T60891	EST	663928
Sequence 13300	T82957	EST	711245
Sequence 13301	T83814	EST	712102

Table 3-5

Sequence 13302	W01580	EST	1273634
Sequence 13303	W04263	EST	1276162
Sequence 13304	W19343	EST	1295012
Sequence 13305	W23474	EST	1300308
Sequence 13306	W69300	EST	1378561
Sequence 13307	W86630	EST	1400227
Sequence 13308	V21239	NUCPATENT	N/A
Sequence 13309	V34331	NUCPATENT	N/A
Sequence 13310	V43615	NUCPATENT	N/A
Sequence 13311	V84583	NUCPATENT	N/A
Sequence 13312	X03841	NUCPATENT	N/A
Sequence 13313	X30393	NUCPATENT	N/A
Sequence 13314	X40045	NUCPATENT	N/A
Sequence 13315	X84983	NUCPATENT	1488593
Sequence 13316	X97975	NUCPATENT	1340080
Sequence 13317	X98010	NUCPATENT	1360094
Sequence 13318	Z14718	NUCPATENT	6383
Sequence 13319	Z40810	NUCPATENT	566550
Sequence 13320	Z41320	NUCPATENT	567459
Sequence 13321	Z41384	NUCPATENT	567528
Sequence 13322	AC02716	PREPATNUC	N/A
Sequence 13323	AC36924	PREPATNUC	N/A
Sequence 13324	AC36925	PREPATNUC	N/A
Sequence 13325	AC36946	PREPATNUC	N/A
Sequence 13326	AC37777	PREPATNUC	N/A
Sequence 13327	AC38546	PREPATNUC	N/A
Sequence 13328	AC39510	PREPATNUC	N/A
Sequence 13329	A02759	Genbank	345130
Sequence 13330	A06846	Genbank	490055
Sequence 13331	A14133	Genbank	490127
Sequence 13332	A16794	Genbank	512417
Sequence 13333	A16796	Genbank	579569
Sequence 13334	A61385	Genbank	3715 <b>797</b>
Sequence 13335	AB000584	Genbank	1813326
Sequence 13336	AB001535	Genbank	3928755
Sequence 13337	AB002303	Genbank	2224550
Sequence 13338	AB002308	Genbank	6634012
Sequence 13339	AB002322	Genbank	6634014
Sequence 13340	AB002323	Genbank	2224590
Sequence 13341	AB002360	Genbank	2224664
Sequence 13342	AB002363	Genbank	2224670
Sequence 13343	AB002364	Genbank	2224672
Sequence 13344 Sequence 13345	AB002366 AB002368	Genbank	2224676
Sequence 13346	AB002387	Genbank	2224680
Sequence 13347		Genbank	2224718
Sequence 13348	AB002391 AB002533	Genbank Genbank	6683696 1944124
Sequence 13349	AB002333	Genbank	
Sequence 13350	AB003723	Genbank	1945608
Sequence 13351	AB003723 AB004064	Genbank	2911141 6983841
Sequence 13352	AB004574	Genbank	3184394
Sequence 13353	AB004857	Genbank	2911111
Sequence 13354	AB004837	Genbank	2924619
Sequence 13355	AB006626	Genbank	6635126
Sequence 13356	AB007144	Genbank	2911155
Sequence 13357	AB007170	Genbank	3077765
40000 1000/	, 1000/1/0	Genbank	0011100

Sequence 13358	AB007874	Genbank	2887448
Sequence 13359	AB007890	Genbank	6634032
Sequence 13360	AB007956	Genbank	3413930
Sequence 13361	AB011083	Genbank	3043545
Sequence 13362	AB011105	Genbank	3043589
Sequence 13363	AB011111	Genbank	3043601
Sequence 13364	AB011120	Genbank	3043619
	AB011148	Genbank	3043675
Sequence 13365	AB011149	Genbank	3043677
Sequence 13366	AB011155	Genbank	3043689
Sequence 13367		Genbank	3327103
Sequence 13368	AB014545	Genbank	3327107
Sequence 13369	AB014547		3327133
Sequence 13370	AB014560	Genbank	3327139
Sequence 13371	AB014563	Genbank	
Sequence 13372	AB014570	Genbank	3327153
Sequence 13373	AB014579	Genbank	3327171
Sequence 13374	AB014589	Genbank	3327191
Sequence 13375	AB014600	Genbank	3327213
Sequence 13376	AB015639	Genbank	5821139
Sequence 13377	AB015752	Genbank	3746106
Sequence 13378	AB017103	Genbank	3493334
Sequence 13379	AB017335	Genbank	3582440
Sequence 13380	AB017548	Genbank	3885363
Sequence 13381	AB017654	Genbank	6979002
Sequence 13382	AB017709	Genbank	3650435
Sequence 13383	AB018257	Genbank	3882148
Sequence 13384	AB018260	Genbank	3882154
Sequence 13385	AB018267	Genbank	3882168
Sequence 13386	AB018319	Genbank	3882272
Sequence 13387	AB018330	Genbank	3882294
Sequence 13388	AB018340	Genbank	3882314
Sequence 13389	AB018342	Genbank	3882318
Sequence 13390	AB018344	Genbank	3882322
Sequence 13391	AB018353	Genbank	3882340
Sequence 13392	AB019038	Genbank	6970469
Sequence 13393	AB019408	Genbank	4587126
Sequence 13394	AB019441	Genbank	4512311
Sequence 13395	AB019524	Genbank	4519939
	AB019527	Genbank	3869126
Sequence 13396		Genbank	3885367
Sequence 13397	AB019564	Genbank	6456471
Sequence 13398	AB020018	Genbank	4760601
Sequence 13399	AB020236	Genbank	6518494
Sequence 13400	AB020335	Genbank	4240148
Sequence 13401	AB020637	<b>C. C</b>	
Sequence 13402	AB020656	Genbank	4240186
Sequence 13403	AB020660	Genbank	4240194
Sequence 13404	AB020662	Genbank	4240198
Sequence 13405	AB020676	Genbank	4240226
Sequence 13406	AB020682	Genbank	4240238
Sequence 13407	AB020694	Genbank	4240262
Sequence 13408	AB020705	Genbank	4240284
Sequence 13409	AB020711	Genbank	4240296
Sequence 13410	AB020859	Genbank	4003379
Sequence 13411	AB020868	Genbank	4003388
Sequence 13412	AB020982	Genbank	4417209
Sequence 13413	AB021470	Genbank	4586521
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<u>Table 3-5</u>

Sequence 13414	AB021663	Genbank	4996450
Sequence 13415	AB022435	Genbank	4996607
Sequence 13416	AB022537	Genbank	6942217
Sequence 13417	AB022785	Genbank	4210446
Sequence 13418	AB023207	Genbank	4589623
Sequence 13419	AB023208	Genbank	4589625
Sequence 13420	AB023224	Genbank	4589657
Sequence 13421	AB023691	Genbank	6519222
Sequence 13422	AB024334	Genbank	6016837
Sequence 13423	AB025355	Genbank	5103146
Sequence 13424	AB025959	Genbank	6520072
Sequence 13425	AB027466	Genbank	6172220
Sequence 13426	AB028624	Genbank	5103045
Sequence 13427	AB028893	Genbank	6552364
Sequence 13428	AB029020	Genbank	5689530
Sequence 13429	AB029331	Genbank	5295999
Sequence 13430	AB030181	Genbank	6730949
Sequence 13431	AB030905	Genbank	5732186
Sequence 13432	AB032951	Genbank	6329748
Sequence 13433	AB032957	Genbank	6329818
Sequence 13434	AB032962	Genbank	6382013
Sequence 13435	AB032969	Genbank	6329965
Sequence 13436	AB033007	Genbank	6330242
Sequence 13437	AB033018	Genbank	6330350
Sequence 13438	AB033020	Genbank	6330364
Sequence 13439	AB033050	Genbank	6330623
Sequence 13440	AB033067	Genbank	6330797
Sequence 13441	AB033069	Genbank	6330811
Sequence 13442	AB033074	Genbank	6330846
Sequence 13443	AB033076	Genbank	6330860
Sequence 13444	AB033079	Genbank	6382025
Sequence 13445	AB033096	Genbank	6331225
Sequence 13446	AB034205	Genbank	6899845
Sequence 13447	AB037763	Genbank	7243064
Sequence 13448	AB037767	Genbank	7243072
Sequence 13449	AB037807	Genbank	7243152
Sequence 13450	AB037813	Genbank	7243164
Sequence 13451	AC000015	Genbank	4263638
Sequence 13452	AC000025	Genbank	4417318
Sequence 13453	AC000046	Genbank	4731059
Sequence 13454	AC000093	Genbank	4678240
Sequence 13455	AC000120	Genbank	1809224
Sequence 13456	AC000353	Genbank	6970735
Sequence 13457	AC000378	Genbank	2270906
Sequence 13458	AC000394	Genbank	2133911
Sequence 13459	AC001226	Genbank	2133862
Sequence 13460	AC001234	Genbank	4028939
Sequence 13461	.AC002038	Genbank	2226439
Sequence 13462	AC002039	Genbank	2342716
Sequence 13463	AC002041	Genbank	2576343
Sequence 13464	- AC002064	Genbank	2076723
Sequence 13465	AC002081	Genbank	2078453
Sequence 13466	AC002094	Genbank	2155224
Sequence 13467	AC002096	Genbank	2599243
Sequence 13468	AC002347	Genbank	2828783
Sequence 13469	AC002350	Genbank	2961443

### <u>Table 3-5</u>

Sequence 13470	AC002365	Genbank	2358015
Sequence 13471	AC002366	Genbank	2739349
Sequence 13472	AC002394	Genbank	2815550
Sequence 13473	AC002395	Genbank	3540146
Sequence 13474	AC002398	Genbank	2529398
Sequence 13475	AC002401	Genbank	2599240
Sequence 13476	AC002487	Genbank	2341013
Sequence 13477	AC002522	Genbank	7124014
Sequence 13478	AC002542	Genbank	2393733
Sequence 13479	AC002544	Genbank	3337382
Sequence 13480	AC002558	Genbank	2580474
Sequence 13481	AC003004	Genbank	2522284
Sequence 13482	AC003007	Genbank	2911728
Sequence 13483	AC003026	Genbank	2815549
Sequence 13484	AC003029	Genbank	9965520
Sequence 13485	AC003037	Genbank	2920805
Sequence 13486	AC003041	Genbank	3264572
Sequence 13487	AC003041	Genbank	10043285
	AC003048	Genbank	2865212
Sequence 13488	AC003070 AC003103	Genbank	2842782
Sequence 13489			3253131
Sequence 13490	AC003104	Genbank	
Sequence 13491	AC003663	Genbank	3097871
Sequence 13492	AC003682	Genbank	3264845
Sequence 13493	AC003688	Genbank	3789719
Sequence 13494	AC003963	Genbank	2995604
Sequence 13495	AC003989	Genbank	2772538
Sequence 13496	AC004006	Genbank	2781390
Sequence 13497	AC004013	Genbank	2781380
Sequence 13498	AC004073	Genbank	2822136
Sequence 13499	AC004084	Genbank	2822156
Sequence 13500	AC004106	Genbank	3150276
Sequence 13501		Genbank	3242744
Sequence 13502	AC004142	Genbank	2880078
Sequence 13503	AC004148	Genbank	3482960
Sequence 13504	AC004152	Genbank	2894631
Sequence 13505	AC004156	Genbank	2896799
Sequence 13506	AC004158	Genbank	2982170
Sequence 13507	AC004221	Genbank	9665053
Sequence 13508	AC004222	Genbank	3108042
Sequence 13509	AC004223	Genbank	3253129
Sequence 13510	AC004227	Genbank	2911718
Sequence 13511	AC004228	Genbank	4263838
Sequence 13512	AC004233	Genbank	2914673
Sequence 13513	AC004236	Genbank	2914668
Sequence 13514	AC004258	Genbank	2924759
Sequence 13515	AC004263	Genbank	2935616
Sequence 13516	AC004381	Genbank	2982169
Sequence 13517	AC004382	Genbank	3252819
Sequence 13518	AC004388	Genbank	3046271
Sequence 13519	AC004408	Genbank	3873185
Sequence 13520	AC004466	Genbank	3617739
Sequence 13521	AC004477	Genbank	3688107
Sequence 13522	AC004494	Genbank	2996647
Sequence 13523	AC004507	Genbank	2996632
Sequence 13524	AC004520	Genbank	3004572
Sequence 13525	AC004522	Genbank	3006227
Dequence 10020	70004752	Germank	JUUULE!

#### <u>Table 3-5</u>

Sequence 13526	AC004527	Genbank	4760422
Sequence 13527	AC004531	Genbank	3337392
Sequence 13528	AC004539	Genbank	3041851
Sequence 13529	AC004582	Genbank	3337307
Sequence 13530	AC004584	Genbank	3417305
Sequence 13531	AC004585	Genbank	3212882
Sequence 13532	AC004587	Genbank	3150014
Sequence 13533	AC004600	Genbank	4388746
Sequence 13534	AC004601	Genbank	3395446
Sequence 13535	AC004611	Genbank	3080643
Sequence 13536	AC004617	Genbank	3738114
Sequence 13537	AC004622	Genbank	3090700
Sequence 13538	AC004634	Genbank	3094993
Sequence 13539	AC004677	Genbank	3126881
Sequence 13540	AC004686	Genbank	3688105
Sequence 13541	AC004760	Genbank	3168626
Sequence 13542	AC004765	Genbank	4731046
Sequence 13543	AC004770	Genbank	3212836
Sequence 13544	AC004798	Genbank	3184497
Sequence 13545	AC004801	Genbank	4204244
Sequence 13546	AC004812	Genbank	3970970
Sequence 13547	AC004813	Genbank	4926912
Sequence 13548	AC004832	Gелралк Селралк	6624129
Sequence 13549	AC004835	Genbank	4508154
Sequence 13550	AC004842	Genbank	4753286
Sequence 13551	AC004846	Genbank	7243869
Sequence 13552	AC004854	Genbank	4827328
Sequence 13553	AC004876	Genbank	4508148
Sequence 13554	AC004890	Genbank	4508146
Sequence 13555	AC004904	Genbank	4156180
Sequence 13556	AC004908	Genbank	4156179
Sequence 13557	AC004938	Genbank	3213059
Sequence 13558	AC004941	Genbank	4454517
Sequence 13559	AC004950	Genbank	4753269
Sequence 13560	AC004955	Genbank	4753267
Sequence 13561	AC004969	Genbank	4156159
Sequence 13562	AC004982	Genbank	3419846
Sequence 13563	AC004983	Genbank	4309885
Sequence 13564	AC004985	Genbank	5708490
Sequence 13565	AC004987	Genbank	3213010
Sequence 13566	AC004990	Genbank	3924668
Sequence 13567	AC004999	Genbank	3970963
Sequence 13568	AC005005	Genbank	4156153
Sequence 13569	AC005007	Genbank	4156151
Sequence 13570	AC005021	Genbank	3694657
Sequence 13571	AC005031	Genbank	3688109
Sequence 13572	AC005037	Genbank	4827310
Sequence 13573	AC005041	Genbank	4508118
Sequence 13574	AC005042	Genbank	4156138
Sequence 13575	AC005046	Genbank	6094632
Sequence 13576	AC005076	Genbank	4508110
Sequence 13577	AC005082	Genbank	5836159
Sequence 13578	AC005084	Genbank	3659503
Sequence 13579	AC005088	Genbank	4753221
Sequence 13580	AC005095	Genbank	4753220
Sequence 13581	AC005104	Genbank	4218027

Sequence 13582	AC005166	Genbank	3242769
Sequence 13583	AC005187	Genbank	3522927
Sequence 13584	AC005189	Genbank	3264580
Sequence 13585	AC005193	Genbank	3309101
Sequence 13586	AC005203	Genbank	3273381
Sequence 13587	AC005212	Genbank	3287444
Sequence 13588	AC005212	Genbank	3287718
Sequence 13589	AC005251	Genbank	3288886
	AC005253 AC005261	Genbank	
Sequence 13590	AC005261 AC005262	Genbank	3289984
Sequence 13591			3289981
Sequence 13592	AC005300	Genbank	6492490
Sequence 13593	AC005318	Genbank	3885345
Sequence 13594	AC005332	Genbank	3659494
Sequence 13595	AC005360	Genbank	3367514
Sequence 13596	AC005363	Genbank	3367511
Sequence 13597	AC005366	Genbank	3367508
Sequence 13598	AC005387	Genbank	3395417
Sequence 13599	AC005391	Genbank	3399673
Sequence 13600	AC005406	Genbank	4580403
Sequence 13601	AC005409	Genbank	4249432
Sequence 13602	AC005410	Genbank	3450753
Sequence 13603	AC005484	Genbank	5091654
Sequence 13604	AC005488	Genbank	5836194
Sequence 13605	AC005509	Genbank	4176355
Sequence 13606	AC005531	Genbank	4153875
Sequence 13607	AC005550	Genbank	3478668
Sequence 13608	AC005562	Genbank	4153858
Sequence 13609	AC005613	Genbank	3540152
Sequence 13610	AC005627	Genbank	7243867
Sequence 13611	AC005629	Genbank	7243877
Sequence 13612	AC005630	Genbank	4159882
Sequence 13613	AC005661	Genbank	3818348
Sequence 13614	AC005668	Genbank	3641771
Sequence 13615	AC005669	Genbank	3702356
Sequence 13616	AC005726	Genbank	3810672
Sequence 13617	AC005730	Genbank	3779042
Sequence 13618	AC005740	Genbank	3687210
Sequence 13619	AC005768	Genbank	6598827
Sequence 13620	AC005785	Genbank	3702290
Sequence 13621	AC005789	Genbank	3702281
Sequence 13622	AC005769	Genbank	3789713
Sequence 13623	AC005829	Genbank	3873300
Sequence 13624	AC005829 AC005837	Genbank	3849820
Sequence 13625	AC005838	Genbank	3947427
Sequence 13626	AC005839	Genbank	4079626
Sequence 13627	AC005841	Genbank	4731044
Sequence 13628	AC005876	Genbank	6249672
Sequence 13629	AC005880	Genbank	6249667
Sequence 13630	AC005899	Genbank	3935221
Sequence 13631	AC005912	Genbank	4165005
Sequence 13632	AC005923	Genbank	4309927
Sequence 13633	AC005924	Genbank	4309926
Sequence 13634	AC005969	Genbank	4309946
Sequence 13635	AC005998	Genbank	5649387
Sequence 13636	AC006009	Genbank	4753278
Sequence 13637	AC006011	Genbank	4508143

Sequence 13638	AC006017	Genbank	4508141
Sequence 13639	AC006023	Genbank	4753260
Sequence 13640	AC006026	Genbank	6042101
Sequence 13641	AC006027	Genbank	4156149
Sequence 13642	AC006050	Genbank	4079627
Sequence 13643	AC006057	Genbank	4731048
Sequence 13644	AC006064	Genbank	4572650
Sequence 13645	AC006084	Genbank	3953485
Sequence 13646	AC006088	Genbank	4204701
Sequence 13647	AC006129	Genbank	3970933
Sequence 13648	AC006137	Genbank	9887710
Sequence 13649	AC006141	Genbank	4580411
Sequence 13650	AC006153	Genbank	4309917
Sequence 13651	AC006160	Genbank	5701616
Sequence 13652	AC006205	Genbank	5815508
Sequence 13653	AC006206	Genbank	4309925
Sequence 13654	AC006207	Genbank	4309924
Sequence 13655	AC006238	Genbank	4204704
Sequence 13656	AC006251	Genbank	4309945
Sequence 13657	AC006254	Genbank	4726099
Sequence 13658	AC006255	Genbank	5306223
Sequence 13659	AC006257	Genbank	4092478
Sequence 13660	AC006273	Genbank	4096060
Sequence 13661	AC006347	Genbank	4309919
Sequence 13662	AC006349	Genbank	7243873
Sequence 13663	AC006409	Genbank	4454486
Sequence 13664	AC006430	Genbank	6453929
Sequence 13665	AC006449	Genbank	6102666
Sequence 13666	AC006450	Genbank	6997336
Sequence 13667	AC006475	Genbank	4753283
Sequence 13668	AC006480	Genbank	6289252
Sequence 13669	AC006487	Genbank	4314418
Sequence 13670	AC006504	Genbank	4220442
Sequence 13671	AC006509	Genbank	5815507
Sequence 13672	AC006512	Genbank	4926863
Sequence 13673	AC006518	Genbank	4713939
Sequence 13674	AC006536	Genbank	4235135
Sequence 13675	AC006539	Genbank	4235141
Sequence 13676	AC006561	Genbank	4558534
Sequence 13677	AC006581	Genbank	4914350
Sequence 13678	AC006597	Genbank	4512725
Sequence 13679 Sequence 13680	AC006930 AC006973	Genbank	4309680
Sequence 13681	AC006973 AC006987	Genbank	4699968
Sequence 13682	AC005967 AC007014	Genbank Genbank	5123989
•	AC007014 AC007029		4371263
Sequence 13683 Sequence 13684	AC007029 AC007038	Genbank	7243872
Sequence 13685	AC007038	Genbank Genbank	5708475 5708471
Sequence 13686	AC007050	Genbank	5708471
Sequence 13687	AC007055	Genbank Genbank	6456174 4885691
Sequence 13688	AC007055 AC007057	Genbank	4680440
Sequence 13689	AC007114	Genbank	4581193
Sequence 13690	AC0071130	Genbank	5001538
Sequence 13691	AC007172	Genbank	4731066
Sequence 13692	AC007172 AC007204	Genbank	4559317
Sequence 13693	AC007207	Genbank	6466489
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### <u>Table 3-5</u>

Sequence 13694	AC007216	Genbank	6806842
Sequence 13695	AC007225	Genbank	6715703
Sequence 13696	AC007229	Genbank	4567173
Sequence 13697	AC007239	Genbank	5523828
Sequence 13698	AC007252	Genbank	5091646
Sequence 13699	AC007314	Genbank	4662678
Sequence 13700	AC007324	Genbank	7923342
Sequence 13701	AC007327	Genbank	4587630
Sequence 13702	AC007390	Genbank	6358845
Sequence 13703	AC007393	Genbank	5306300
Sequence 13704	AC007446	Genbank	5069495
Sequence 13705	AC007528	Genbank	5263309
Sequence 13706	AC007536	Genbank	5917929
Sequence 13707	AC007541	Genbank	4982536
Sequence 13708	AC007546	Genbank	5668756
Sequence 13709	AC007551	Genbank	4809347
Sequence 13710	AC007566	Genbank	4835815
Sequence 13711	AC007766	Genbank	5030437
Sequence 13712	AC007848	Genbank	5923645
Sequence 13712	AC007860	Genbank	5649179
Sequence 13714	AC007868	Genbank	6041758
Sequence 13715	AC007934	Genbank	6721208
Sequence 13716	AC008009	Genbank	5668757
Sequence 13717	AC008018	Genbank	7958973
Sequence 13718	AC008102	Genbank	9797815
Sequence 13719	AC008115	Genbank	5801654
Sequence 13720	AC008123	Genbank	6006046
Sequence 13721	AC008151	Genbank	5629925
Sequence 13722	AC008498	Genbank	6850299
Sequence 13723	AC008545	Genbank	6721145
Sequence 13724	AC008583	Genbank	7158894
Sequence 13725	AC008893	Genbank	7259691
Sequence 13726	AC008981	Genbank	6721142
Sequence 13727	AC009298	Genbank	7243882
Sequence 13728	AC009363	Genbank	6648145
Sequence 13729	AC009509	Genbank	6492473
Sequence 13730	AC009946	Genbank	6604542
Sequence 13731	AC010077	Genbank	5870275
Seguence 13732	AC010168	Genbank	6855156
Sequence 13733	AC010197	Genbank	6143841
Sequence 13734	AC010436	Genbank	6938861
Sequence 13735	AC010525	Genbank	7248932
Sequence 13736	AC010582	Genbank	6721135
Sequence 13737	AC010722	Genbank	6587939
Sequence 13738	AC011331	Genbank	9929686
Sequence 13739	AC011599	Genbank	7113871
Sequence 13740	AC012085	Genbank	6634755
Sequence 13741	AC012087	Genbank	7114465
Sequence 13742	AC016831	Genbank	6539289
Sequence 13743	AC016941	Genbank	8567654
Sequence 13744	AC020663	Genbank	6682593
Sequence 13745	AC021049	Genbank	7248920
Sequence 13746	AF000982	Genbank	2580549
Sequence 13747	AF001549	Genbank	3355302
Sequence 13748	AF001552	Genbank	2339976
Sequence 13749	AF002668	Genbank	2232173

Page 66 of 105

#### <u>Table 3-5</u>

Sequence 13750	AF004162	Genbank	3046385
Sequence 13751	AF004337	Genbank	2245557
Sequence 13752	AF004338	Genbank	2245558
Sequence 13753	AF004339	Genbank	2245559
Sequence 13754	AF004342	Genbank	2245565
Sequence 13755	AF006513	Genbank	2645428
Sequence 13756	AF006621	Genbank	7629276
Sequence 13757	AF007134	Genbank	2852608
Sequence 13758	AF007146	Genbank	2852624
Sequence 13759	AF007189	Genbank	2459927
Sequence 13760	AF007189	Genbank	
,			2970122
Sequence 13761	AF007872	Genbank	2358280
Sequence 13762	AF007875	Genbank	2258417
Sequence 13763	AF008303	Genbank	2580619
Sequence 13764	AF009242	Genbank	2338289
Sequence 13765	AF010313	Genbank	6468761
Sequence 13766	AF011889	Genbank	2335186
Sequence 13767	AF013488	Genbank	2343184
Sequence 13768	AF013758	Genbank	3046899
Sequence 13769	AF013970	Genbank	2801421
Sequence 13770	AF014398	Genbank	8148061
Sequence 13771	AF015812	Genbank	2599359
Sequence 13772	AF017790	Genbank	2501872
Sequence 13773	AF020352	Genbank	2655054
Sequence 13774	AF020782	Genbank	2522480
Sequence 13775	AF021819	Genbank	2460317
Sequence 13776	AF022211	Genbank	2465723
Sequence 13777	AF022728	Genbank	2935182
Sequence 13778	AF026802	Genbank	5712085
Sequence 13779	AF026852	Genbank	3599965
Sequence 13780	AF027205	Genbank	2598967
Sequence 13781	AF028832	Genbank	3287488
Sequence 13782	AF029750	Genbank	2587057
Sequence 13783	AF029786	Genbank	3403166
Sequence 13784	AF031379	Genbank	4894208
Sequence 13785	AF035286	Genbank	2661038
Sequence 13786	AF035296	Genbank	2661052
Sequence 13787	AF035304	Genbank	2661064
Sequence 13788	AF035840	Genbank	3800739
Sequence 13789	AF036715	Genbank	4007385
Sequence 13790	AF036718	Genbank	2708629
Sequence 13791	AF037447	Genbank	6466790
Sequence 13792	AF038200	Genbank	2795921
Sequence 13793	AF038406	Genbank	3523122
Sequence 13794	AF039019	Genbank	2828109
Sequence 13795	AF039575	Genbank	2773157
Sequence 13796	AF039594	Genbank	3366667
Sequence 13797	AF040628	Genbank	3869269
Sequence 13798	AF042836	Genbank	3776241
Sequence 13799	AF042636 AF043105	Genbank Genbank	3776241 3169541
		Genbank Genbank	
Sequence 13800	AF044956		5326827
Sequence 13801	AF044957	Genbank	4164445
Sequence 13802	AF044959	Genbank	3348136
Sequence 13803		Genbank	2865655
Sequence 13804	AF047184	Genbank	2909859
Sequence 13805	AF047185	Genbank	2909861

Page 67 of 105

# Table 3-5

AF047438	Genbank	3335131
· · · · ·	Genbank	3335135
		2921872
		5326819
		4164453
		3360398
	= :	3360438
	T	3360475
		3360473
		3510694
•		2967847
	<b>4.0</b>	
		3135305
		4092053
		3005703
		3273727
		nil
		3649607
		3063652
		4165042
		3047377
AF058988		3300089
AF059527		4454661
AF060798	Genbank	3372665
AF061737	Genbank	4335938
AF062077	Genbank	3641524
AF064019	Genbank	3347856
AF064861	Genbank	3171154
AF067139	Genbank	3337440
AF068302	Genbank	4584876
AF068754	Genbank	3283408
AF068755	Genbank	4321979
AF069378	Genbank	3982733
AF069532	Genbank	3892190
AF070523	Genbank	3764088
AF070539	Genbank	3387896
AF070556	Genbank	3387921
AF070630	Genbank	3283898
AF070635	Genbank	3283905
AF070638	Genbank	3283909
AF070646	Genbank	3283920
AF070649	Genbank	3283923
AF070655	Genbank	4454685
AF070659	Genbank	4454693
AF070669	Genbank	4454713
AF070673	Genbank	3978241
	Genbank	5725511
		3641537
AF075061	Genbank	3377602
	Genbank	4689107
AF077030	Genbank Genbank	4689107 4689119
AF077030 AF077036	Genbank	4689119
AF077030 AF077036 AF077043	Genbank Genbank	4689119 4689133
AF077030 AF077036 AF077043 AF077044	Genbank Genbank Genbank	4689119 4689133 4689135
AF077030 AF077036 AF077043 AF077044 AF077052	Genbank Genbank Genbank Genbank	4689119 4689133 4689135 4689151
AF077030 AF077036 AF077043 AF077044 AF077052 AF077200	Genbank Genbank Genbank Genbank Genbank	4689119 4689133 4689135 4689151 4679013
AF077030 AF077036 AF077043 AF077044 AF077052	Genbank Genbank Genbank Genbank	4689119 4689133 4689135 4689151
	AF060798 AF061737 AF062077 AF064019 AF064861 AF067139 AF068302 AF068754 AF068755 AF069378 AF069532 AF070523 AF070523 AF070539 AF070630 AF070635 AF070638 AF070646 AF070649 AF070655 AF070659	AF047440 Genbank AF047472 Genbank AF050638 Genbank AF050639 Genbank AF052092 Genbank AF052129 Genbank AF052182 Genbank AF052573 Genbank AF052578 Genbank AF053356 Genbank AF054990 Genbank AF055474 Genbank AF055584 Genbank AF057356 Genbank AF057356 Genbank AF057357 Genbank AF057357 Genbank AF05735 Genbank AF05735 Genbank AF05735 Genbank AF05735 Genbank AF05735 Genbank AF05735 Genbank AF058988 Genbank AF058988 Genbank AF068757 Genbank AF061737 Genbank AF061737 Genbank AF064019 Genbank AF064019 Genbank AF064019 Genbank AF064861 Genbank AF068754 Genbank AF068755 Genbank AF068755 Genbank AF0689532 Genbank AF070659 Genbank AF070638 Genbank AF070659 Genbank AF070649 Genbank AF070659 Genbank AF070669 Genbank AF070669 Genbank AF070669 Genbank AF070673 Genbank AF070673 Genbank

Page 68 of 105

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**Table 3-5** 

Sequence 13862	AF078850	Genbank	5531814
Sequence 13863	AF082283	Genbank	4092066
Sequence 13864	AF082513	Genbank	3649656
Sequence 13865	AF083441	Genbank	5813822
Sequence 13866	AF085844	Genbank	3483158
Sequence 13867	AF086210	Genbank	3483555
•			
Sequence 13868	AF086329	Genbank	3483674
Sequence 13869	AF086337	Genbank	3483682
Sequence 13870	AF086378	Genbank	3483723
Sequence 13871	AF086490	Genbank	3483835
Sequence 13872	AF086557	Genbank	3483902
Sequence 13873	AF086564	Genbank	3483909
Sequence 13874	AF086628	Genbank	4240457
Sequence 13875	AF087020	Genbank	3851144
Sequence 13876	AF087160	Genbank	3869380
Sequence 13877	AF087959	Genbank	3523165
Sequence 13878	AF088060	Genbank	3523266
Sequence 13879	AF088071	Genbank	3523277
Sequence 13880	AF090095	Genbank	4063630
Sequence 13881	AF090947	Genbank	6690255
Sequence 13882	AF091075	Genbank	3859987
Sequence 13883	AF092094	Genbank	4426606
Sequence 13884	AF092124	Genbank	3659900
Sequence 13885	AF092128	Genbank	5138905
Sequence 13886	AF092130	Genbank	5138909
Sequence 13887	AF092135	Genbank	5138919
Sequence 13888	AF097026	Genbank	6409120
Sequence 13889	AF097514	Genbank	4808600
Sequence 13890	AF098162	Genbank	3929582
Sequence 13891	AF100615	Genbank	4808624
Sequence 13892	AF100616	Genbank	4808626
Sequence 13893	AF100018 AF100748	Genbank	5410281
Sequence 13894	AF100757	Genbank	5410299
Sequence 13895	AF100760	Genbank	5410305
Sequence 13896	AF102803	Genbank	4092759
Sequence 13897	AF104233	Genbank	5880481
Sequence 13898	AF104913	Genbank	3941723
Sequence 13899	AF106622	Genbank	4378528
Sequence 13900	AF106681	Genbank	5410327
Sequence 13901	AF109134	Genbank	4139227
Sequence 13902	AF109718	Genbank	3978547
Sequence 13903	AF110460	Genbank	4324698
Sequence 13904	AF111847	Genbank	7211441
Sequence 13905	AF112213	Genbank	6563213
Sequence 13906	AF112222	Genbank	6563 <b>229</b>
Sequence 13907	AF112227	Genbank	4545218
Sequence 13908	AF113011	Genbank	6642745
Sequence 13909	AF113016	Genbank	6642755
Sequence 13910	AF113141	Genbank	4512031
Sequence 13911	AF113535	Genbank	6523824
Sequence 13912	AF113540	Genbank	6523834
Sequence 13913	AF113615	Genbank	5106955
Sequence 13914	AF113680	Genbank	6855605
Sequence 13915	AF113690	Genbank	6855619
Sequence 13916	AF113700	Genbank	6855634
Sequence 13917	AF113701	Genbank	6855635
•			<del>-</del>

**Table 3-5** 

Sequence 13918	AF115402	Genbank	4559272
Sequence 13919	AF116910	Genbank	4768837
Sequence 13920	AF117235	Genbank	6563243
Sequence 13921	AF117888	Genbank	5732617
Sequence 13922	AF118084	Genbank	6650813
		Genbank	4235636
Sequence 13923	AF118124		
Sequence 13924	AF118649	Genbank	6840946
Sequence 13925	AF119297	Genbank	4633508
Sequence 13926	AF123052	Genbank	5106556
Sequence 13927	AF125530	Genbank	6563259
Sequence 13928	AF126023	Genbank	6563275
Sequence 13929	AF126782	Genbank	6318547
Sequence 13930	AF127577	Genbank	7019596
Sequence 13931	AF128527	Genbank	4928043
Sequence 13932	AF128536	Genbank	5305705
Sequence 13933	AF129756	Genbank	4337095
Sequence 13934	AF131738	Genbank	4406548
Sequence 13935	AF131748	Genbank	4406563
Sequence 13936	AF131758	Genbank	4406576
Sequence 13937	AF131771	Genbank	4406595
•	AF131777	Genbank	4406602
Sequence 13938			
Sequence 13939	AF131781	Genbank	4406608
Sequence 13940	AF131814	Genbank	4406648
Sequence 13941	AF131838	Genbank	4406677
Sequence 13942	AF131856	Genbank	4406702
Sequence 13943	AF132939	Genbank	4680648
Sequence 13944	AF132942	Genbank	4680654
Sequence 13945	AF132973	Genbank	4680716
Sequence 13946	AF132984	Genbank	4959567
Sequence 13947	AF134726	Genbank	4529886
Sequence 13948	AF135162	Genbank	7259481
Sequence 13949	AF135488	Genbank	5668619
Sequence 13950	AF146651	Genbank	5020073
Sequence 13951	AF147330	Genbank	4761681
Sequence 13952	AF147378	Genbank	<b>4</b> 761 <b>729</b>
Sequence 13953	AF147412	Genbank .	4761763
Sequence 13954	AF150089	Genbank	5107165
Sequence 13955	AF151052	Genbank	7106825
Sequence 13956	AF151103	Genbank	5758136
Sequence 13957	AF151109	Genbank	6166337
Sequence 13958	AF151801	Genbank	4929554
Sequence 13959	AF151807	Genbank	4929566
Sequence 13960	AF151832	Genbank	4929616
Sequence 13961	AF151835	Genbank	4929622
Sequence 13962	AF151878	Genbank	4929708
Sequence 13963	AF151903	Genbank	4929758
Sequence 13964	AF151905	Genbank	4929762
Sequence 13965	AF151906	Genbank	4929764
Sequence 13966	AF152363	Genbank	5669134
Sequence 13967	AF153273	Genbank	5932030
Sequence 13968	AF153274	Genbank	5932030 5932031
Sequence 13969	AF153274 AF153608	Genbank	
Sequence 13970	AF156965		5231140
	AF158965 AF158180	Genbank	5731112
Sequence 13971		Genbank	6959307
Sequence 13972	AF159295	Genbank	5714635
Sequence 13973	AF161384	Genbank	6841181

Sequence 13974	AF161389	Genbank	6841191
Sequence 13975	AF161428	Genbank	6841269
Sequence 13976	AF161448	Genbank	6841309
Sequence 13977	AF161464	Genbank	6841451
Sequence 13978	AF161479	Genbank	6841481
Sequence 13979	AF161481	Genbank	6841485
Sequence 13980	AF161485	Genbank	6841493
Sequence 13981	AF161500	Genbank	6841523
Sequence 13982	AF161503	Genbank	6841529
Sequence 13983	AF161509	Genbank	6841541
Sequence 13984	AF161512	Genbank	6841547
Sequence 13985	AF161541	Genbank	6841349
Sequence 13986	AF161556	Genbank	6841379
Sequence 13987	AF165926	Genbank	5817858
	AF168956	Genbank	5702387
Sequence 13988		Genbank	
Sequence 13989	AF172066		5733811
Sequence 13990	AF176315	Genbank	5823349
Sequence 13991	AF181896	Genbank	6272683
Sequence 13992	AF187320	Genbank	6164847
Sequence 13993	AF187554	Genbank	6653225
Sequence 13994	AF188745	Genbank	6425043
Sequence 13995	AF189712	Genbank	6760474
Sequence 13996	AF190465	Genbank	6120105
Sequence 13997	AF191018	Genbank	6457339
Sequence 13998	AF191298	Genbank	7656642
Sequence 13999	AF201077	Genbank	6456748
Sequence 14000	AF203815	Genbank	6979641
Sequence 14001	AF205588	Genbank	6531675
Sequence 14002	AF207550	Genbank	6470333
Sequence 14003	AF216305	Genbank	6942010
Sequence 14004	AF228703	Genbank	7158868
Sequence 14005	AF228704	Genbank	715 <b>887</b> 1
Sequence 14006	AJ000519	Genbank	2739214
Sequence 14007	AJ001014	Genbank	3171909
Sequence 14008	AJ001866	Genbank	2969902
Sequence 14009	AJ001981	Genbank	2546978
Sequence 14010	AJ002030	Genbank	2570006
Sequence 14011	AJ002308	Genbank	2959871
Sequence 14012	AJ002385	Genbank	2597930
Sequence 14013	AJ006470	Genbank	3687321
Sequence 14014	AJ010069	Genbank	3483012
Sequence 14015	AJ010597	` Genbank	3559873
Sequence 14016	AJ012409	Genbank	3881975
Sequence 14017	AJ012499	Genbank	5441359
Sequence 14018	AJ223812	Genbank	2894518
Sequence 14019	AJ224442	Genbank	2911586
Sequence 14020	AJ230595	Genbank	4583774
Sequence 14021	AJ230596	Genbank	4583775
Sequence 14022	AJ230605	Genbank	4583784
Sequence 14023	AJ238243	Genbank	4826530
Sequence 14024	AJ23B283	Genbank	4704219
Sequence 14025	AJ238374	Genbank	6634418
Sequence 14026	AJ245719	Genbank	6688168
Sequence 14027	AJ249731	Genbank	5921469
Sequence 14028	AJ250042	Genbank	6013005
Sequence 14029	AJ250915	Genbank	6996445
20400100 140E9	, 10200010	Company	

Sequence 14030	AJ251961	Genbank	6634453
Sequence 14031	AJ271216	Genbank	6966966
Sequence 14032	AK000018	Genbank	7019828
Sequence 14033	AK000023	Genbank	7019837
Sequence 14034	AK000035	Genbank	7019855
Sequence 14035	AK000208	Genbank	7020141
Sequence 14036	AK000291	Genbank	7020274
Sequence 14037	AK000348	Genbank	7020373
Sequence 14038	AK000345	Genbank	7020402
Sequence 14039	AK000395	Genbank	7020402
Sequence 14040	AK000393 AK000426	Genbank	7020455
Sequence 14041	AK000428	Genbank	
Sequence 14042	AK000454	Genbank	7020520
Sequence 14043	AK000455 AK000462	Genbank	7020550
Sequence 14043	AK000462 AK000470		7020566
		Genbank	7020580
Sequence 14045	AK000507	Genbank	7020644
Sequence 14046	AK000548	Genbank	7020717
Sequence 14047	AK000560	Genbank	7020738
Sequence 14048	AK000571	Genbank	7020756
Sequence 14049	AK000585	Genbank	7020779
Sequence 14050	AK000615	Genbank	702 <b>0827</b>
Sequence 14051	AK000620	Genbank	7020834
Sequence 14052	AK000624	Genbank	7020839
Sequence 14053	AK000655	Genbank	7020888
Sequence 14054	AK000667	Genbank	7020906
Sequence 14055	AK000669	Genbank	7020909
Sequence 14056	AK000721	Genbank	7020985
Sequence 14057	AK000725	Genbank	7020990
Sequence 14058	AK000739	Genbank	7021014
Sequence 14059	AK000745	Genbank	7021025
Sequence 14060	AK000752	Genbank	7021035
Sequence 14061	AK000765	Genbank	7021058
Sequence 14062	AK000800	Genbank	7021104
Sequence 14063	AK000818	Genbank	7021128
Sequence 14064	AK000826	Genbank	7021139
Sequence 14065	AK000866	Genbank	7021190
Sequence 14066	AK000891	Genbank	7021840
Sequence 14067	AK000933	Genbank	7021908
Sequence 14068	AK000934	Genbank	7021909
Sequence 14069	AK000959	Genbank	7021945
Sequence 14070	AK000991	Genbank	7021994
Sequence 14071	AK001006	Genbank	70 <b>22017</b>
Sequence 14072	AK001093	Genbank	7022142
Sequence 14073	AK001105	Genbank	7022160
Sequence 14074	AK001154	Genbank	7022228
Sequence 14075	AK001237	Genbank	702 <b>2366</b>
Sequence 14076	AK001285	Genbank	702 <b>2444</b>
Sequence 14077	AK001330	Genbank	7022520
Sequence 14078	AK001387	Genbank	7022615
Sequence 14079	AK001413	Genbank	7022655
Sequence 14080	AK001441	Genbank -	7022699
Sequence 14081	AK001451	Gen <b>ban</b> k	70 <b>22717</b>
Sequence 14082	AK001469	Genbank	702 <b>2747</b>
Sequence 14083	AK001493	Genba <b>n</b> k	702 <b>2783</b>
Sequence 14084	AK001536	Genbank	7022851
Sequence 14085	AK001571	Genbank	7022906

Sequence 14086	AK001656	Genbank	7023047
Sequence 14087	AK001690	Genbank	7023105
Sequence 14088	AK001691	Genbank	7023106
Sequence 14089	AK001717	Genbank	7023151
Sequence 14090	AK001718	Genbank	7023153
Sequence 14091	AK001747	Genbank	7023203
Sequence 14092	AK001762	Genbank	7023231
Sequence 14093	AK001855	Genbank	7023231
Sequence 14094	AK001877	Genbank	7023362
Sequence 14095	AK001878	Genbank	7023418
Sequence 14096	AK001961	Genbank	
Sequence 14097	AK002002	Genbank	7023549
Sequence 14098	AK002002 AK002046	Genbank	7023620
Sequence 14099	AK002048 AK002102		7023689
Sequence 14100	AK002102 AK002169	Genbank	7023781
•		Genbank	7023884
Sequence 14101	AK002190	Genbank	7023914
Sequence 14102	AL008582	Genbank	5360981
Sequence 14103	AL008639	Genbank	6599068
Sequence 14104	AL008721	Genbank	3171883
Sequence 14105	AL008725	Genbank	2791551
Sequence 14106	AL009051	Genbank	2995195
Sequence 14107	AL009181	Genbank	2853179
Sequence 14108	AL020993	Genbank	3980107
Sequence 14109	AL020995	Genbank	5881341
Sequence 14110	AL021069	Genbank	2853183
Sequence 14111	AL021368	Genbank	3080468
Sequence 14112	AL021546	Genbank	28268 <b>90</b>
Sequence 14113	AL021920	Genbank	3264535
Sequence 14114	AL022165	Genbank	3281985
Sequence 14115	AL022238	Genbank	4176442
Sequence 14116	AL022240	Genbank	4826471
Sequence 14117	AL022302	Genbank	5262844
Sequence 14118	AL022312	Genbank	4914501
Sequence 14119	AL022313	Genbank	42003 <b>26</b>
Sequence 14120	AL022316	Genbank	4691242
Sequence 14121	AL022320	Genbank	4914513
Sequence 14122	AL022328	Genbank	526301 <b>0</b>
Sequence 14123	AL022394	Genbank	95 <b>81758</b>
Sequence 14124	AL022395	Genbank	44682 <b>87</b>
Sequence 14125	AL022725	Genbank	5679748
Sequence 14126	AL023284	Genbank	3355875
Sequence 14127	AL023494	Genbank	5360992
Sequence 14128	AL023798	Genbank	3550031
Sequence 14129	AL024493	Genbank	328843 <b>9</b>
Sequence 14130	AL030996	Genbank	36 <b>88349</b>
Sequence 14131	AL031003	Genbank	4007185
Sequence 14132	AL031259	Genbank	3790132
Sequence 14133	AL031274	Genbank	342108 <b>3</b>
Sequence 14134	AL031287	Genbank	4 <b>914533</b>
Sequence 14135	AL031295	Genbank	4376011
Sequence 14136	AL031368	Genbank	4455593
Sequence 14137	AL031390	Genbank	5002609
Sequence 14138	AL031428	Genbank	59 <b>31</b> 719
Sequence 14139	AL031431	Genbank	4938290
Sequence 14140	AL031432	Genbank	4375969
Sequence 14141	AL031433	Genbank	4826442

Sequence 14142	AL031447	Genbank	4826431
Sequence 14143	AL031597	Genbank	4775640
Sequence 14144	AL031662	Genbank	9716901
Sequence 14145	AL031667	Genbank	9408727
Sequence 14146	AL031681	Genbank	9408729
Sequence 14147	AL031710	Genbank	9798435
Sequence 14148	AL031716	Genbank	4826474
Sequence 14149	AL031732	Genbank	4581343
Sequence 14150	AL031781	Genbank	4038570
Sequence 14151	AL032821	Genbank	4572584
Sequence 14152	AL033377	Genbank	4826462
Sequence 14153	AL033397	Genbank	4902626
Sequence 14154	AL033522	Genbank	4191258
Sequence 14155	AL033531	Genbank	6807582
Sequence 14156	AL033543	Genbank	4581346
Sequence 14157	AL034417	Genbank	5102616
Sequence 14158	AL034548	Genbank	7263904
Sequence 14159	AL035209	Genbank	4160217
Sequence 14160	AL035413	Genbank	6010110
Sequence 14161	AL035413 AL035420	Genbank	
Sequence 14162	AL035425	Genbank	5918161
Sequence 14163	AL035458	Genbank	6465823
Sequence 14164	AL035456 AL035541		6624641
Sequence 14165	AL035652	Genbank Genbank	9581764
Sequence 14166	AL035681		9650702
Sequence 14167	AL035699	Genbank	4902689
Sequence 14167	AL033699 AL049249	Genbank	4826515
Sequence 14169	AL049249 AL049282	Genbank	4499987
Sequence 14170	AL049202 AL049404	Genbank	4500041
Sequence 14171	AL049404 AL049557	Genbank Genbank	4500192
Sequence 14172	AL049563	_	55 <b>96765</b>
Sequence 14173		Genbank	4902753
Sequence 14174	AL049564 AL049589	Genbank Genbank	4902757
Sequence 14175	AL049569 AL049696		56 <b>79567</b>
Sequence 14176	AL049090 AL049742	Genbank Genbank	5931893
Sequence 14177	AL049742 AL049761		5419783
Sequence 14178	AL049766	Genbank	5738437
Sequence 14179	AL049766 AL049776	Genbank	5763746
Sequence 14180	AL049779	Genbank	6681704
Sequence 14181	AL049779 AL049829	Genbank Genbank	8176894
Sequence 14182	AL049829 AL049844	Genbank	8217859
Sequence 14183	AL049869	Genbank	5924019
Sequence 14184	AL049809 AL049929		8217861
Sequence 14185	AL049968	Genbank	4884174
Sequence 14186	AL049987	Genbank	4884217
Sequence 14187		Genbank	4884238
Sequence 14188	AL050021 AL050027	Genbank	4884264
		Genbank	4884266
Sequence 14189 Sequence 14190	AL050035	Genbank	4884276
Sequence 14190 Sequence 14191	AL050083	Genbank Canbank	4884317
Sequence 14191 Sequence 14192	AL050161	Genbank	4884375
_ •	AL050170	Genbank	4884165
Sequence 14193	AL050215	Genbank	4884454
Sequence 14194	AL050273	Genbank	4886450
Sequence 14195	AL050275	Genbank	4886500
Sequence 14196	AL050290	Genbank	4886512
Sequence 14197	AL050343	Genbank	61 <b>37008</b>

<u>Table 3-5</u>

Sequence 14198	AL050363	Genbank	4914597
Sequence 14199	AL050383	Genbank	4914585
Sequence 14200	AL050392	Genbank	4914613
Sequence 14201	AL078591	Genbank	6562085
Sequence 14202	AL078621	Genbank	6013067
Sequence 14203	AL079295	Genbank	5102607
Sequence 14204	AL079304	Genbank	8176897
Sequence 14205	AL079342	Genbank	6018784
Sequence 14206	AL080088	Genbank	5262504
Sequence 14207	AL080089	Genbank	5262504
Sequence 14208	AL080097	Genbank	5262519
Sequence 14209	AL080109	Genbank	5262535
Sequence 14210	AL080135	Genbank	5262576
Sequence 14211	AL080144	Genbank	5262592
Sequence 14212	AL080186	Genbank	5262664
Sequence 14213	AL080241	Genbank	5730204
Sequence 14214	AL080250	Genbank	5804917
Sequence 14215	AL080276	Genbank	5763753
Sequence 14216	AL096773	Genbank	
Sequence 14217	AL096803	Genbank	5918011
Sequence 14218	AL109657	Genbank	6010162
Sequence 14219	AL109684	Genbank	6136991
Sequence 14220	AL109884 AL109925	Genbank	5689805
Sequence 14221	AL110183	Genbank	7105935
Sequence 14222	AL110185		5817095
Sequence 14223	AL110193	Genbank	5817098
Sequence 14224	AL110193	Genbank Genbank	5817109
Sequence 14225	AL110202	Genbank	5817111
Sequence 14226	AL110202 AL110208	Genbank	5817121 5817127
Sequence 14227	AL110262	Genbank	5817127 5817224
Sequence 14228	AL117334	Genbank	7248316
Sequence 14229	AL117424	Genbank	5911856
Sequence 14230	AL117430	Genbank	5911865
Sequence 14231	AL117442	Genbank	5911885
Sequence 14232	AL117558	Genbank	5912097
Sequence 14233	AL117599	Genbank	5912167
Sequence 14234	AL117621	Genbank	5912202
Sequence 14235	AL117648	Genbank	5912240
Sequence 14236	AL117662	Genbank	5912257
Sequence 14237	AL117666	Genbank	5912264
Sequence 14238	AL121576	Genbank	8176910
Sequence 14239	AL121580	Genbank	6456854
Sequence 14240	AL121603	Genbank	6434634
Sequence 14241	AL121656	Genbank	7159617
Sequence 14242	AL121782	Genbank	6456831
Sequence 14243	AL121790	Genbank	8919824
Sequence 14244	AL121913	Genbank	7161781
Sequence 14245	AL121963	Genbank	7161783
Sequence 14246	AL122003	Genbank	6580480
Sequence 14247	AL122013	Genbank	7242500
Sequence 14248	AL122050	Genbank	6093244
Sequence 14249	AL122066	Genbank	6102860
Sequence 14250	AL122089	Genbank	6102899
Sequence 14251	AL133051	Genbank	6453468
Sequence 14252	AL133060	Genbank	6453483
Sequence 14253	AL133094	Genbank	6453548

# <u>Table 3-5</u>

Sequence 14254	AL133238	Genbank	6624577
Sequence 14255	AL133241	Genbank	7630038
Sequence 14256	AL133243	Genbank	6491713
Sequence 14257	AL133246	Genbank	7594587
Sequence 14258	AL133370	Genbank	8248722
Sequence 14259	AL133453	Genbank	7009594
Sequence 14260	AL133455	Genbank	7019734
Sequence 14261	AL133555	Genbank	6599122
Sequence 14262	AL135744	Genbank	6682291
Sequence 14263	AL135858	Genbank	7159622
Sequence 14264	AL136296	Genbank	8217913
Sequence 14265	AL136504	Genbank	7018294
Sequence 14266	AL137440	Genbank	6808001
Sequence 14267	AL137450	Genbank	6808024
Sequence 14268	AL137489	Genbank	
	AL137496	Genbank	6808110
Sequence 14269			6808137
Sequence 14270	AL137553	Genbank	6808238
Sequence 14271	AL137567	Genbank	6808276
Sequence 14272	AL137659	Genbank	6807778
Sequence 14273	AL137673	Genbank	6807841
Sequence 14274	AL137681	Genbank	6807931
Sequence 14275	AL137714	Genbank	6808128
Sequence 14276	AL137750	Genbank	6808386
Sequence 14277	AL157428	Genbank	7018458
Sequence 14278	AL157464	Genbank	7018479
Sequence 14279	AP000008	Genbank	3132318
Sequence 14280	AP000031	Genbank	3132341
Sequence 14281	AP000038	Genbank	3132348
Sequence 14282	AP000351	Genbank	6069325
Sequence 14283	AP000354	Genbank	5103017
Sequence 14284	AP000473	Genbank	6942329
Sequence 14285	AP000525	Genbank	59315 <b>03</b>
Sequence 14286	AP000526	Genbank	5931504
Sequence 14287	AP000962	Genbank	6942330
Sequence 14288	AP001055	Genbank	6693605
Sequence 14289	AP001136	Genbank.	6970360
Sequence 14290	AP001137	Genbank	6970361
Sequence 14291	AP001172	Genbank	6983884
Sequence 14292	AP001343	Genbank	7209831
Sequence 14293	AP001346	Genbank	7209834
Sequence 14294	D10493	Genbank	219932
Sequence 14295	D12485	Genbank	219943
Sequence 14296	D14082	Genbank	468933
Sequence 14297	D14533	Genbank	286028
Sequence 14298	D14540	Genbank	467511
Sequence 14299	D14658	Genbank	285940
Sequence 14300	D14665	Genbank	6630617
Sequence 14301	D14696	Genbank	285962
Sequence 14302	D14710	Genbank	559324
Sequence 14303	D21089	Genbank	475156
Sequence 14304	D21235	Genbank	498145
Sequence 14305	D21262	Genbank	434764
Sequence 14306	D25215	Genbank	517114
Sequence 14307	D25274	Genbank	464185
Sequence 14308	D26600	Genbank	565650
Sequence 14309	D28468	Genbank	460704
		Company	700704

<u>Table 3-5</u>

Sequence 14310	D29011	Genbank	558525
Sequence 14311	D29643	Genbank	473936
Sequence 14312	D29805	Genbank	474986
Sequence 14313	D29954	Genbank	473940
Sequence 14314	D29958	Genbank	473948
Sequence 14315	D30648	Genbank	506337
Sequence 14316	D31885	Genbank	505097
Sequence 14317	D32053	Genbank	2366751
Sequence 14318	· D38112	Genbank	644480
Sequence 14319	D38524	Genbank	633070
Sequence 14320	D42041	Genbank	577294
Sequence 14321	D42063	Genbank	924266
Seguence 14322	D49489	Genbank	1136742
Sequence 14323	D50420	Genbank	2618577
Sequence 14324	D50525	Genbank	1167502
Sequence 14325	D50916	Genbank	1469174
Sequence 14326	D50929	Genbank	1469200
Sequence 14327	D55673	Genbank	870746
Sequence 14328	D63486	Genbank	1469885
Sequence 14329	D63880	Genbank	961451
Sequence 14330	D64015	Genbank	2281005
Sequence 14331	D79205	Genbank	1754620
Sequence 14332	D79986	Genbank	1136389
Sequence 14333	D80000	Genbank	1136415
Sequence 14334	D83004	Genbank	1181557
Sequence 14335	D84273	Genbank	3894217
Sequence 14336	D86962	Genbank	1503997
Sequence 14337	D86985	Genbank	6634002
Sequence 14338	D87003	Genbank	2114242
Sequence 14339	D87127	Genbank	1817551
Sequence 14340	D87328	Genbank	1817331
Sequence 14341	D87666	Genbank	1620016
Sequence 14342	D87684	Genbank	1663703
Sequence 14343	D90209	Genbank	220087
Sequence 14344	E00882	Genbank	2169143
Sequence 14345	E01094	Genbank	2169353
Sequence 14346	E01533	Genbank	2169789
Sequence 14347	E01932	Genbank	2170180
Sequence 14348	E01954	Genbank	2170202
Sequence 14349	E01979	Genbank	2170227
Sequence 14350	E02164	Genbank	2170402
Sequence 14351	E02628	Genbank	2170856
Sequence 14352	E05647	Genbank	2173834
Sequence 14353	E05957	Genbank	2174144
Sequence 14354	E08663	Genbank	2176776
Sequence 14355	E12274	Genbank	3251108
Sequence 14356	E12457	Genbank	3251290
Sequence 14357	E12651	Genbank	3251483
Sequence 14358	E13330	Genbank	3252135
Sequence 14359	E13405	Genbank	3252210
Sequence 14360	J00139	Genbank	182715
Sequence 14361	J01415	Genbank	1944628
Sequence 14362	J02642	Genbank	182862
Sequence 14363	J02960	Genbank	178203
Sequence 14364	J03248	Genbank	183053
Sequence 14365	J03459	Genbank	187172
	000-00	Sombank	10/1/2

Page 77 of 105

**Table 3-5** 

Sequence 14366	J03537	Genbank	337513
Sequence 14367	J03592	Genbank	339722
Sequence 14368	J03746	Genbank	183655
Sequence 14369	J03779	Genbank	179833
Sequence 14370	J03934	Genbank	189245
Sequence 14371	J04173	Genbank	551173
Sequence 14372		Genbank	190371
Sequence 14373		Genbank	189425
Sequence 14374		Genbank	186504
Sequence 14375		Genbank	340020
Sequence 14376		Genbank	337377
Sequence 14377		Genbank	184097
Sequence 14378		Genbank	292462
Sequence 14379		Genbank	307365
Sequence 14380		Genbank	388033
Sequence 14381	L05491	Genbank	292831
Sequence 14382		Genbank	1835658
Sequence 14383		Genbank	292442
Sequence 14384	L07287	Genbank	292434
Sequence 14385		Genbank '	
Sequence 14386		Genbank	184250
Sequence 14387		Genbank	179295
Sequence 14388	L09054	Genbank	307344
Sequence 14389		Genbank	186522
Sequence 14390	L10612	Genbank	402701
Sequence 14391	L11066	Genbank	307322
Sequence 14392	L11370	Genbank	387674
Sequence 14393	L12136	Genbank	181536
Sequence 14394	L12387	Genbank	459835
Sequence 14395	L12711	Genbank	388890
Sequence 14396	L13278	Genbank	292414
Sequence 14397	L14778	Genbank	306476
Sequence 14398		Genbank	440307
Sequence 14399	L19437	Genbank	4995970
Sequence 14400	L19739	Genbank	431318
Sequence 14401	L19783	Genbank	404725
Sequence 14402	L20046	Genbank	306741
Sequence 14403	L20941	Genbank	507251
Sequence 14404	L22569	Genbank	348706
Sequence 14405	L24444	Genbank	495860
Sequence 14406	L25931	Genbank	438638
Sequence 14407	L28809	Genbank	454151
Sequence 14408 Sequence 14409	L28997 L29766	Genbank Genbank	607027
Sequence 14410	L31610	Genbank	537525 1220360
Sequence 14410	L37127		4164098
Sequence 14411	L37127 L38995	Genbank Genbank	704415
Sequence 14413	L39320	Genbank	1226044
Sequence 14414	L39326	Genbank	1228094
Sequence 14415	L40396	Genbank	887357
Sequence 14416	L43575	Genbank	899064
Sequence 14417	L43964	Genbank	951202
Sequence 14418	L44140	Genbank	1203968
Sequence 14419	L47162	Genbank	1082035
Sequence 14420	L47647	Genbank	1002033
Sequence 14421	L76159	Genbank	1246232
	=:		

Table 3-5

Sequence 14422	L76416	Genbank	5566605
Sequence 14423	M10036	Genbank	339840
Sequence 14424	M11146	Genbank	182504
Sequence 14425	M11147	Genbank	18251 <b>3</b>
Sequence 14426	M11353	Genbank	184092
Sequence 14427	M12623	Genbank	184233
Sequence 14428	M15205	Genbank	339718
Sequence 14429	M15796	Genbank	181271
Sequence 14430	M16541	Genbank	1280204
Sequence 14431	M17886	Genbank	190233
Sequence 14432	M18048	Genbank	337699
Sequence 14433	M19997	Genbank	181968
Sequence 14434	M21142	Genbank	183402
Sequence 14435	M21575	Genbank	1311702
Sequence 14436	M21895	Genbank	189523
Sequence 14437	M22760	Genbank	695359
Sequence 14438	M22918	Genbank	189019
Sequence 14439	M22920	Genbank	189021
Sequence 14440	M23410	Genbank	762884
Sequence 14441	M24070	Genbank	181485
Sequence 14442	M24194	Genbank	187701
Sequence 14443	M24486	Genbank	190785
Sequence 14444	M24543	Genbank	341200
Sequence 14445	M24690	Genbank	341218
Sequence 14446	M26481	Genbank	619789
Sequence 14447	M26663	Genbank	618463
Sequence 14448	M29064	Genbank	337452
Sequence 14449	M29960	Genbank	339886
Sequence 14450	M31468	Genbank	190876
Sequence 14451	M31520	Genbank	337504
Sequence 14452	M36341	Genbank	178984
Sequence 14453	M36981	Genbank	189239
Sequence 14454	M37104	Genbank	179274
Sequence 14455	M37583	Genbank	184059
Sequence 14456	M38188	Genbank	189378
Sequence 14457	M55150	Genbank	182392
Sequence 14458	M55409	Genbank	189596
Sequence 14459	M58485	Genbank	180154
Sequence 14460	M60457	Genbank	181249
Sequence 14461	M60484	Genbank	190225
Sequence 14462	M60756	Genbank	184085
Sequence 14463	M60857	Genbank	181334
Sequence 14464	M61764	Genbank	183702
Sequence 14465	M63180	Genbank	339679
Sequence 14466	M73547	Genbank	190161
Sequence 14467	M74775	Genbank	187151
Sequence 14468	M75099	Genbank	337369
Sequence 14469	M76299	Genbank	188778
Sequence 14470	M77232	Genbank	307392
Sequence 14471	M86181	Genbank	190566
Sequence 14472	M87339	Genbank	1498255
Sequence 14473	M88006	Genbank	178522
Sequence 14474 Sequence 14475	M88468	Genbank Genbank	307197 170577
Sequence 14475	M90357 M93036	Genbank Genbank	179577
Sequence 14476 Sequence 14477	M93651		182904
Cedaelice 144//	14192021	Genbank	338038

Page 79 of 105

Sequence 14478	M94556	Genbank	188855
Sequence 14479	M96982	Genbank	338262
Sequence 14480	S48220	Genbank	257451
Sequence 14481	S54761	Genbank	265221
Sequence 14482	S72871	Genbank	639594
Sequence 14483	S73498	Genbank	688010
Sequence 14484	S73591	Genbank	688296
Sequence 14485	S74678	Genbank	241477
Sequence 14486	S81752	Genbank	1438780
Sequence 14487	S82597	Genbank	1699343
Sequence 14488	S82616	Genbank	1699307
Sequence 14489	U0023B	Genbank	404860
Sequence 14490	U02031	Genbank	451329
Sequence 14491	U03886	Genbank	458225
Sequence 14492	U07231	Genbank	517195
Sequence 14493	U07857	Genbank	469048
Sequence 14494	U09466	Genbank	495492
Sequence 14495	U09813	Genbank	1008454
Sequence 14496	U10248	Genbank	984280
Sequence 14497	U10362	Genbank	505651
Sequence 14498	U12404	Genbank	
Sequence 14499	U12465	Genbank	531170
Sequence 14500	U12897		562073
Sequence 14501	U13616	Genbank Genbank	562075
Sequence 14502	U14193	Genbank	608024
Sequence 14503	U14970	Genbank	555899
Sequence 14504	U14972	Genbank	550020
Sequence 14505	U14973	Genbank	550024 550026
Sequence 14506	U17899	Genbank	
Sequence 14507	U18321	Genbank	717053
Sequence 14508	U18671	Genbank	603763
Sequence 14509	U20285	Genbank	1293919
Sequence 14510	U20362	Genbank	644878
Sequence 14511	U25182	Genbank	755485
Sequence 14512	U29171	Genbank	799380 881618
Sequence 14513	U29589	Genbank	
Sequence 14514	U30826	Genbank	903978 1 <b>0</b> 49079
Sequence 14515	U34683	Genbank	1236349
Sequence 14516	U39817	Genbank	
Sequence 14517	U41060	Genbank	1072121 1256000
Sequence 14518	U41195	Genbank	1244694
Sequence 14519	U41448	Genbank	3335023
Sequence 14520	U41668	Genbank	1477481
Sequence 14521	U44798	Genbank	1174216
Sequence 14522	U46751		
Sequence 14523	U49245	Genbank Genbank	3077821 1216503
Sequence 14524	U51559	Genbank	
Sequence 14525	U52111	Genbank	1236709
Sequence 14526	U52852	_	8331754
Sequence 14527	U54993	Genbank Genbank	8370402
Sequence 14528	U56833	Genbank Genbank	4097314
Sequence 14529			1465750
	U57846	Genbank	1373418
Sequence 14530	U57847	Genbank	1373420
Sequence 14531 Sequence 14532	U59911	Genbank	1403712
	U66243	Genbank	1772645
Sequence 14533	U66589	Genbank	<b>157</b> 5566

<u>Table 3-5</u>

Sequence 14534	U67243	Genbank	2228516
Sequence 14535	U68566	Genbank	1916621
Sequence 14536	U68758	Genbank	4097815
Sequence 14537	U72648	Genbank	3914602
Sequence 14538	U73168	Genbank	1613899
Sequence 14539	U73645	Genbank	1737200
Sequence 14540	U73824	Genbank	1857236
Sequence 14541	U75285	Genbank	2315862
Sequence 14542	U76549	Gen <b>bank</b>	1673574
Sequence 14543	U76669	Gen <b>bank</b>	6672211
Sequence 14544	U77643	Genbank	2062390
Sequence 14545	U80017	Genbank	1737211
Sequence 14546	U80460	Genbank	2341006
Sequence 14547	U85625	Genbank	5091494
Sequence 14548	U87460	Genbank	2076881
Sequence 14549	U90904	Genbank	1913882
Sequence 14550	U90907	Genbank	1913886
Sequence 14551	U91327	Genbank	
Sequence 14552	U91328	Genbank	1871209
•	U94586		2088550
Sequence 14553	U95689	Genbank Genbank	1946691
Sequence 14554			2004000
Sequence 14555	U96629	Genbank	2341008
Sequence 14556 Sequence 14557	V00478	Genbank	28244
	V00662	Genbank	13003
Sequence 14558	X00351	Genbank	28251
Sequence 14559	X02152	Genbank	34312
Sequence 14560	X03083	Genbank	34309
Sequence 14561	X03205	Genbank	36162
Sequence 14562	X05276	Genbank	37201
Sequence 14563	X05332	Genbank	35740
Sequence 14564	X05857	Genbank	31988
Sequence 14565	X06409	Genbank	30204
Sequence 14566	X14608	Genbank	296365
Sequence 14567	X15822	Genbank	30146
Sequence 14568	X15940	Genbank	36129
Sequence 14569	X16135	Genbank	32355
Sequence 14570	X16560	Genbank	30154
Sequence 14571	X17025	Genbank	488749
Sequence 14572	X17620	Genbank	35067
Sequence 14573	X52839	Genbank	36125
Sequence 14574	X52851	Genbank	30167
Sequence 14575	X52882	Genbank	311380
Sequence 14576	X52897	Genbank	24484
Sequence 14577	X52943	Genbank	28912
Sequence 14578	X56932	Genbank	23690
Sequence 14579	X56998	Genbank	37564
Sequence 14580	X58288	Genbank	32455
Sequence 14581	X59434	Genbank	432375
Sequence 14582	X60221	Genbank	509290
Sequence 14583	X61755	Genbank	32387
Sequence 14584	X63432	Genbank	28335
Sequence 14585	X63657	Genbank	296185
Sequence 14586	X64123	Genbank	35819
Sequence 14587	X65923	Genbank	31302
Sequence 14588	X66975	Gen <b>bank</b>	32353
Sequence 14589	X67247	Genbank	36149

Sequence 14590	X67698	Genbank	37476
Sequence 14591	X69910	Genbank	297407
Sequence 14592	X70940	Genbank	38455
Sequence 14593	X71428	Genbank	393415
Sequence 14594	X72841	Genbank	297903
Sequence 14595	X74801	Genbank	671526
Sequence 14596	X74979	Genbank	400462
Sequence 14597	X75593	Genbank	452319
Sequence 14598	X75684	Genbank	429092
Sequence 14599	X75861	Genbank	456258
Sequence 14600	X76013	Genbank	531595
Sequence 14601	X78137	Genbank	460770
Sequence 14602	X80200	Genbank	951276
Sequence 14603	X81713	Genbank	558213
Sequence 14604	X81788	Genbank	1045058
Sequence 14605	X81895	Genbank	2370158
Sequence 14606	X81900	Genbank	2274973
Sequence 14607	X84908	Genbank	1502344
Sequence 14608	X85372	Genbank	806563
Sequence 14609	X87241	Genbank	1107686
Sequence 14610	X87344	Genbank	1054740
Sequence 14611	X87949	Genbank	1143491
Sequence 14612	X91249	Genbank	1160185
Sequence 14613	X93036	Genbank	1085025
Sequence 14614	X94910	<b>G</b> enbank	3413292
Sequence 14615	X95592	Genbank	1185118
Sequence 14616	X96586	Genbank	1556398
Sequence 14617	X97544	Genbank	1770563
Sequence 14618	X98356	Genba <b>nk</b>	1648869
Sequence 14619	X99585	Genba <b>n</b> k	1770518
Sequence 14620	Y00052	Genbank	30308
Sequence 14621	Y00282	Genbank	36048
Sequence 14622	Y00345	Genbank	35569
Sequence 14623 Sequence 14624	Y00815	Genbank	34266
Sequence 14625	Y08201 Y09836	Genbank	1552548
Sequence 14626	Y10345	Genbank Genbank	1743253
Sequence 14627	Y15286	Genbank	2292905
Sequence 14628	Z11692	Genbank	2584788 31107
Sequence 14629	Z11793	Genbank	36425
Sequence 14630	Z13009	Genbank	31072
Sequence 14631	Z14244	Genbank	30150
Sequence 14632	Z21507	Genbank	38521
Sequence 14633	Z24724	Genbank	505034
Sequence 14634	Z25749	Genbank	550116
Sequence 14635	Z31695	Genbank	469143
Sequence 14636	Z36832	Genbank	533946
Sequence 14637	Z36849	Genbank	533963
Sequence 14638	Z46973	Genbank	987947
Sequence 14639	Z47087	Genbank	860989
Sequence 14640	Z48950	Genbank	761715
Sequence 14641	Z54246	Genbank	1000700
Sequence 14642	Z68164	Genbank	1107692
Sequence 14643	<b>Z68284</b>	Genbank	1130698
Sequence 14644	Z69711	Genbank	1204109
Sequence 14645	Z69890	<b>G</b> enba <b>nk</b>	1213361

Sequence 14646	Z73900	Genbank	1340069
Sequence 14647	Z82242	Genbank	1841915
Sequence 14648	Z82975	Genbank	1673497
Sequence 14649	Z84474	Genbank	1903188
Sequence 14650	Z84812	Genbank	1845164
Sequence 14651	Z85986	Genbank	4034056
Sequence 14652	Z92544	Genbank	1869775
Sequence 14653	Z92845	Genbank	2462406
Sequence 14654	Z93041	Genbank	1905788
	Z93928	Genbank	2578051
Sequence 14655		Genbank	6572188
Sequence 14656	Z95113	Genbank	2342581
Sequence 14657	Z95126	Genbank	2121307
Sequence 14658	Z95704	*·-··-	
Sequence 14659	Z97053	Genbank	9650676
Sequence 14660	Z97054	Genbank	2370077
Sequence 14661	Z97056	Genbank	2832593
Sequence 14662	Z97353	Genbank	4455632
Sequence 14663	Z97652	Genbank	6456822
Sequence 14664	Z98049	Genbank	2462401
Sequence 14665	Z98749	Genbank	4775611
Sequence 14666	Z98884	Genbank	5304861
Sequence 14667	Z98950	Genbank	<b>30</b> 367 <b>83</b>
Sequence 14668	Z99128	Genbank	3135974
Sequence 14669	Z99916	Genbank	4158122
Sequence 14670	AA004759	EST	1448327
Sequence 14671	AA007319	EST	14634 <b>64</b>
Sequence 14672	AA009421	EST	1470619
Sequence 14673	AA009598	EST	1470757
Sequence 14674	AA010071	EST	1471099
Sequence 14675	AA010798	EST	1471844
Sequence 14676	AA011369	EST	1472466
Sequence 14677	AA011539	EST	1472565
Sequence 14678	AA011569	EST	1472595
Sequence 14679	AA013065	EST	1474101
Sequence 14680	AA018720	EST	1481994
Sequence 14681	AA019036	EST	1482627
Sequence 14682	AA019504	EST	1482187
Sequence 14683	AA020851	EST	1484622
Sequence 14684	AA021306	EST	1484995
Sequence 14685	AA024597	EST	1489502
Sequence 14686	AA025057	EST	1489962
Sequence 14687	AA025122	EST	<b>1</b> 490 <b>03</b> 7
Sequence 14688	AA026215	EST	1492901
Sequence 14689	AA026316	EST	1492253
Sequence 14690	AA026353	EST	1492674
Sequence 14691	AA026643	EST	1492801
Sequence 14692	AA026758	EST	1492556
Sequence 14693	AA027229	EST	1492149
Sequence 14694	AA029679	· EST	1497082
Sequence 14695	AA034414	EST	1506241
Sequence 14696	AA035260	EST	1506772
Sequence 14697	AA035377	EST	1507101
Sequence 14698	AA037143	EST	1512251
Sequence 14699	AA039589	EST	1516014
Sequence 14700	AA039683	EST	1516155
Sequence 14701	AA039734	EST	1516236
25425.55 .5.51			

**Table 3-5** 

Sequence 14702	AA039876	EST	1516172
Sequence 14703	AA039929	EST	1516206
Sequence 14704	AA040404	EST	1516765
Sequence 14705	AA043048	EST	1522583
Sequence 14706	AA043193	EST	1521111
Sequence 14707	AA043606	EST	1521472
Sequence 14708	AA044663	EST	1523043
Sequence 14709	AA044835	EST	1523038
Sequence 14710	AA046542	EST	1524439
Sequence 14711	AA046810	EST	1524915
Sequence 14712	AA046864	EST	1524763
Sequence 14713	AA047046	EST	1524944
Sequence 14714	AA053607	EST	1544534
Sequence 14715	AA053853	EST	1544797
Sequence 14716'	AA053926	EST	1544870
Sequence 14717	AA053942	EST	1545004
Sequence 14718	AA054471	EST	1545396
Sequence 14719	AA056107	EST	1548444
Sequence 14720	AA057352	EST	1549991
Sequence 14721	AA057541	EST	1550246
Sequence 14722	AA057559	EST	1550199
Sequence 14723	AA063079	EST	1556632
Sequence 14724	AA063612	EST	1557579
Sequence 14725	AA065328	EST	1929208
Sequence 14726	AA069023	EST	1576401
Sequence 14727	AA069422	EST	1576781
Sequence 14728	AA069655	EST	1577014
Sequence 14729	AA070277	EST	1577637
Sequence 14730	AA070855	EST	1578487
Sequence 14731	AA071269	EST	1578757
Sequence 14732	AA071402	EST	1578773
Sequence 14733	AA071472	EST	1578903
Sequence 14734	AA074221	EST	1614090
Sequence 14735	AA074611	EST	1614677
Sequence 14736	AA074614	EST	1614492
Sequence 14737	AA074664	EST	1614608
Sequence 14738	AA074831	EST	1614875
Sequence 14739	AA075200	EST	1615070
Sequence 14740	AA075981	EST	1615850
Sequence 14741	AA079050	EST	1617942
Sequence 14742	AA081478	EST	1623536
Sequence 14743	AA082545	EST	1624602
Sequence 14744	AA083712	EST	1625772
Sequence 14745	AA084678	EST	1626735
Sequence 14746	AA084868	EST	1626988
Sequence 14747	AA088189	EST	1633764
Sequence 14748	AA090105	EST	1636589
Sequence 14749	AA092310	EST	1637531
Sequence 14750	AA092324	EST	1637545
Sequence 14751	AA098978	EST	1644862
Sequence 14752	AA099015	EST	1644895
Sequence 14753	AA099080	EST	1645522
Sequence 14754	AA099327	EST	1646033
Sequence 14755	AA100639	EST	1646940
Sequence 14756	AA101116	EST	1647692
Sequence 14757	AA101598	EST	1648552
•		· <del>- •</del>	
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<u>Table 3-5</u>

Sequence	14758	AA101663	EST	1648744
Sequence 1		AA102442	EST	1647356
Sequence		AA102560	EST	1647752
Sequence		AA114042	EST	1667919
Sequence		AA115474	EST	1670145
Sequence :		AA126478	EST	1686071
Sequence :		AA126762	EST	1686262
Sequence :		AA126841	EST	1686323
Sequence :	14766	AA127483	EST	1686755
Sequence :	14767	AA127550	EST	1686822
Sequence 1	14768	AA127886	EST	1687193
Sequence 1	14769	AA129205	EST	1688990
Sequence :		AA129465	EST	1689336
Sequence 1		AA129935	EST	1689511
Sequence 1		AA131336	EST	1692834
Sequence		AA132089	EST	1693579
Sequence 1		AA132394	EST	
Sequence 1		AA132989		1693885
			EST	1694505
Sequence 1		AA133031	EST	1694539
Sequence 1		AA133736	EST	1690768
Sequence 1		AA133759	EST	1690727
Sequence 1		AA134263	EST	1691637
Sequence 1		AA134609	EST	1695597
Sequence 1	14781	AA135030	EST	- 1696141
Sequence 1		AA135126	EST	1696227
Sequence 1	4783	AA135254	EST	1696373
Sequence 1	14784	AA136337	EST	1697545
Sequence 1	4785	AA136723	EST	1697933
Sequence 1		AA136761	EST	1698008
Sequence 1		AA137016	EST	1698225
Sequence 1		AA142942	EST	1712320
Sequence 1		AA143324	EST	1712695
Sequence 1		AA143422	EST	1712852
Sequence 1		AA147059	EST	1716433
Sequence 1		AA147948	EST	1717455
Sequence 1		AA148889	EST	1718998
Sequence 1		AA149404	EST	
Sequence 1		AA149518	EST	1719920
Sequence 1				1720336
Sequence 1		AA149579	EST	1720380
		AA149801	EST	1720899
Sequence 1		AA149826	EST	1720961
Sequence 1		AA150361	EST	1721873
Sequence 1		AA150481	EST	1721995
Sequence 1		AA151140	EST	1719383
Sequence 1		AA151423	EST	1719820
Sequence 1		AA151795	EST	1720490
Sequence 1		AA151809	EST	1720504
Sequence 1	4805	AA151816	EST	1720511
Sequence 1	4806	AA152179	EST	1721231
Sequence 1	4807	AA155999	EST	1727666
Sequence 1	4808	AA156226	EST	1727844
Sequence 1		AA158153	EST	1732990
Sequence 1		AA158669	EST	1733480
Sequence 1	-	AA158704	EST	1733533
Sequence 1		AA159176	EST	1733969
Sequence 1		AA159347	EST	1733909
Coquonioe 1	7010	, v110004/	E31	1734149

<u>Table 3-5</u>

Sequence 148	314 AA160	0666	EST	1736043
Sequence 148	315 AA161		EST	1735906
Sequence 148			EST ·	1741453
Sequence 148			EST	1745155
Sequence 148				1745024
Sequence 148		_	EST	1745322
Sequence 148				1745386
Sequence 148				1745886
Sequence 148				1748121
Sequence 148				1750791
Sequence 148				1753550
Sequence 148				1753883
Sequence 148		100		
Sequence 148				1754068 1757375
Sequence 148				1757713
Sequence 148				1760308
Sequence 148				1760572
Sequence 148		-		1765368
Sequence 148		_ : _ :		1766581
Sequence 148				1766108
Sequence 148				1774446
Sequence 148				1775103
Sequence 148				1779239
Sequence 148				1781923
Sequence 148				1781585
Sequence 148				1784213
Sequence 148			_	1 <b>78</b> 4783
Sequence 148				1799204
Sequence 148				1802881
Sequence 148		-		1801341
Sequence 148				1801826
Sequence 148				1802148
Sequence 148		_		1809555
Sequence 148				1813173
Sequence 148				1 <b>8</b> 15385
Sequence 148		047 E	EST ·	1816018
Sequence 148		-	EST ·	1816048
Sequence 148			EST ·	1833202
Sequence 148		941 E	EST :	1839687
Sequence 1485			EST 1	1844515
Sequence 148			EST 1	1846276
Sequence 1489			EST 1	1846765
Sequence 148		485 E	EST 1	1846812
Sequence 1489			EST 1	1846970
Sequence 1489		B14 E	EST 1	1847200
Sequence 148	59 AA226	768 E		1848466
Sequence 1486	60 AA227	285 E	ST 1	848839
Sequence 1486	31 AA229			1851876
Sequence 1486	62 AA232			855538
Sequence 1486	33 AA235		_	860273
Sequence 1486				874187
Sequence 1486				874456
Sequence 1486				874594
Sequence 1486				875185
Sequence 1486				879338
Sequence 1486				886090
				200030

<u>Table 3-5</u>

Sequence	14870	AA252998	EST	1882921
Sequence		AA257006	EST	1891135
Sequence		AA258030	EST	1894480
Sequence		AA258271	EST	1893453
Sequence		AA258394	EST	1893536
Sequence		AA261894	EST	1897728
Sequence		AA261960	EST	1898149
Sequence		AA262181	EST	1898452
Sequence	14878	AA262425	EST	1897921
Sequence	14879	AA262871	EST	1898592
Sequence	14880	AA263047	EST	1898845
Sequence	14881	AA278243	EST	1920183
Sequence		AA278787	EST	1920108
Sequence		AA279139	EST	1920605
Sequence		AA279421	EST	1920886
Sequence		AA280206	EST	1921881
Sequence		AA280658	EST	1923453
Sequence		AA280848	EST	1923546
		AA282253		
Sequence :			EST	1925187
Sequence 1		AA285040	EST	1927721
Sequence :	14890	AA285126	EST	1928107
Sequence '		AA292069	EST	1940055
Sequence		AA292296	EST	1940276
Sequence 1		AA294863	EST	1947208
Sequence 1		AA295158	EST	1947513
Sequence :		AA296277	EST	1948652
Sequence:		AA296690	EST	1949182
Sequence 1		AA297999	EST	1950582
Sequence:		AA300796	EST	1953128
Sequence 1	14899	AA303856	EST	1956421
Sequence 1		AA304244	EST	1956597
Sequence 1	14901	AA304272	EST	1956615
Sequence:	14902	AA305667	EST	1958242
Sequence :	14903	AA306121	EST	1958449
Sequence 1	14904	AA306470	EST	1958798
Sequence 1	14905	AA307154	EST	1959482
Sequence 1	14906	AA307818	EST	1960145
Sequence 1		AA308297	EST	1960626
Sequence :		AA308596	EST	1961163
Sequence 1		AA311749	EST	1964076
Sequence 1		AA313317	EST	1965646
Sequence 1		AA313802	EST	1966131
Sequence 1		AA314008	EST	1966337
Sequence 1		AA314493	EST	1966822
Sequence 1		AA314869	EST	1967218
Sequence 1		AA315072	EST	1967411
Sequence 1		AA315192		
Sequence 1		AA316110	EST	1967541
			EST	1968439
Sequence 1		AA316272	EST	1968601
Sequence 1		AA316868	EST	1969196
Sequence 1		AA317537	EST	1969886
Sequence 1		AA318361	EST	1970709
Sequence 1		AA319855	EST	1972181
Sequence 1		AA325081	EST	1977579
Sequence 1		AA328973	EST	1981216
Sequence 1	14925	AA329553	EST	1982040

**Table 3-5** 

Sequence 14926	AA334936	EST	1987254
Sequence 14927	AA339220	EST	1991456
Sequence 14928	AA341254	EST	1993491
Sequence 14929	AA342547	EST	1994804
Sequence 14930	AA343888	EST	1996146
Sequence 14931	AA348129	EST	2000440
Sequence 14932	AA355326	EST	2007708
Sequence 14933	AA359995	EST	2012346
Sequence 14934	AA362311	EST	2014781
Sequence 14935	AA362547	EST	2014761
Sequence 14936	AA363030	EST	2014918
Sequence 14937	AA363971	EST	2016289
Sequence 14938	AA364506	EST	2016845
Sequence 14939	AA371595	EST	2024137
Sequence 14940	AA371677	EST	2024220
Sequence 14941	AA371861	EST	2024387
Sequence 14942	AA376859	EST	2029176
Sequence 14943	AA377534	EST	2029863
Sequence 14944	AA378981	EST	2031371
Sequence 14945	AA379089	EST	2031419
Sequence 14946	AA382161	EST	2034479
Sequence 14947	AA382953	EST	2035271
Sequence 14948	AA393055	EST	2046151
Sequence 14949	AA393236	EST	2046205
Sequence 14950	AA397442	EST	2050483
Sequence 14951	AA398254	EST	2051363
Sequence 14952	AA400319	EST	2054395
Sequence 14953	AA400551	EST	2054422
Sequence 14954	AA400632	EST	2054503
Sequence 14955	AA401792	EST	2057276
Sequence 14956	AA402613	EST	2056402
Sequence 14957	AA402800	EST	2056553
Sequence 14958	AA404535	EST	2059268
Sequence 14959	AA404961	EST	2063336
Sequence 14960	AA405814	EST	2063797
Sequence 14961	AA406106	EST	2064105
Sequence 14962	AA406543	EST	2064657
Sequence 14963	AA410626	EST	2069851
Sequence 14964	AA411013	EST	2070137
Sequence 14965	AA411736	EST	2069397
Sequence 14966	AA412286	EST	2070857
Sequence 14967 Sequence 14968	AA412695 AA417129	EST	2071301
Sequence 14969	AA417129 AA417185	EST EST	2077272
Sequence 14970	AA418317	EST	2077292
Sequence 14971	AA418533	EST	2080146
Sequence 14972	AA418734	EST	2080314
Sequence 14973	AA419146	EST	2080535 2078944
Sequence 14974	AA420564	EST	2076944
Sequence 14975	AA420675	EST	2094553
Sequence 14976	AA420705	EST	2094553
Sequence 14977	AA420873	EST	2094306
Sequence 14978	AA421675	EST	2100533
Sequence 14979	AA422114	EST	2100938
Sequence 14980	AA424037	EST	2103938
Sequence 14981	AA425108	EST	2107178

Sequerice 14982	AA425723	EST	2106443
Sequence 14983	AA426125	EST	2106597
Sequence 14984	AA426618	EST	2107298
Sequence 14985	AA427438	EST	2111308
Sequence 14986	AA427994	EST	2112068
Sequence 14987	AA427999	EST	2112073
Sequence 14988	AA428399	EST	2110228
Sequence 14989	AA431300	EST	2115008
	AA431320	· EST	2115028
Sequence 14990	AA431322	EST	2115030
Sequence 14991	AA431581	EST	2115289
Sequence 14992		EST	2115520
Sequence 14993	AA431812	EST	2114670
Sequence 14994	AA432282		2139054
Sequence 14995	AA434140	EST	
Sequence 14996	AA436116	EST	2141030
Sequence 14997	AA436991	EST	2141905
Sequence 14998	AA437166	EST	2142080
Sequence 14999	AA437224	EST	2142138
Sequence 15000	AA442133	EST	2154011
Sequence 15001	AA443297	EST	2155972
Sequence 15002	AA443762	EST	2156437
Sequence 15003	AA446569	EST	2159234
Sequence 15004	AA447974	EST	2161 <del>644</del>
Sequence 15005	AA448292	EST	2161962
Sequence 15006	AA451954	EST	2165 <b>623</b>
Sequence 15007	AA453183	EST	2166852
Sequence 15008	AA456223	EST	2179433
Sequence 15009	AA456863	EST	2179583
Sequence 15010	AA458476	EST	2183383
Sequence 15011	AA459991	EST	2184875
Sequence 15012	AA461239	EST	2186359
Sequence 15013	AA461254	EST	2186374
Sequence 15014	AA463447	EST	2188331
Sequence 15015	AA463899	EST	218878 <b>3</b>
Sequence 15016	AA464099	EST	2188983
Sequence 15017	AA464353	EST	2189237
Sequence 15018	AA465191	EST	2191358
Sequence 15019	AA465537	EST	2191704
Sequence 15020	AA468049	EST	2194583
Sequence 15021	AA468360	EST	2194894
Sequence 15022	AA470144	EST	2197453
Sequence 15023	AA477119	EST	2205803
Sequence 15024	AA478414	EST	2207048
Sequence 15025	AA478637	EST	2207271
Sequence 15026	AA479488	EST	2208044
Sequence 15027	AA479863	EST	2205749
Sequence 15028	AA479892	EST	2204374
Sequence 15029	AA481045	EST	2210597
Sequence 15030	AA483162	EST	2211975
Sequence 15031	AA483450	EST	2212263
Sequence 15032	AA484155	EST	2212968
	AA487382	EST	2217546
Sequence 15033	AA487628	EST	2217792
Sequence 15034			2217835
Sequence 15035	AA487671	EST	
Sequence 15036	AA488200	EST	2215631
Sequence 15037	AA488782	EST	2218384

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# Table 3-5

PCT/US01/02318

Sequence 15038	AA492042	EST	2221604
Sequence 15039	AA492143	EST	2221705
Sequence 15040	AA492267	EST	2221829
Sequence 15041	AA492423	EST	2221985
Sequence 15042	AA493364	EST	2223205
Sequence 15043	AA493382	EST	2223223
Sequence 15044	AA493522	EST	2223363
Sequence 15045	AA493624	EST	2223465
Sequence 15046	AA493807	EST	2223648
Sequence 15047	AA496002	EST	2229323
Sequence 15048	AA502198	EST	2237165
Sequence 15049	AA503141	EST	2238108
Sequence 15050	AA503404	EST	2238371
Sequence 15051	AA503941	EST	2238908
Sequence 15052	AA504095	EST	2240255
Sequence 15053	AA504531	EST	2240691
Sequence 15054	AA504703	EST	2240863
Sequence 15055	AA505440	EST	2241577
Sequence 15056	AA506953	EST	
Sequence 15057	AA508597	EST	2243392
Sequence 15058	AA513357	EST	2246100
Sequence 15059	AA513367	EST	2251769 2251779
Sequence 15060	AA514331	EST	2253839
Sequence 15061	AA514474	EST	2254074
Sequence 15062	AA514538	EST	2254138
Sequence 15063	AA515736	EST	2255336
Sequence 15064	AA520996	EST	2261539
Sequence 15065	AA521142	EST	2261685
Sequence 15066	AA521353	EST	2261896
Sequence 15067	AA522584	EST	2263296
Sequence 15068	AA523005	EST	2388772
Sequence 15069	AA524825	EST	2265753
Sequence 15070	AA525374	EST	2266302
Sequence 15071	AA525960	EST	2268029
Sequence 15072	AA526186	EST	2268255
Sequence 15073	AA526243	EST	2268312
Sequence 15074	AA526569	EST	2268638
Sequence 15075	AA527150	EST	2269219
Sequence 15076	AA527943	EST	2270012
Sequence 15077	AA528066	EST	2270135
Sequence 15078	AA531276	EST	2273982
Sequence 15079	AA531295	EST	2274001
Sequence 15080	AA532807	EST	2278383
Sequence 15081	AA533483	EST	2277579
Sequence 15082	AA534137	EST	2278153
Sequence 15083	AA534340	EST	2278593
Sequence 15084	AA535571	EST	2279824
Sequence 15085	AA535832	EST	2280085
Sequence 15086	AA541343	EST	2287777
Sequence 15087	AA541537	EST	2287971
Sequence 15088	AA541666	EST	2288100
Sequence 15089	AA548238	EST	2318520
Sequence 15090	AA551141	EST	2321393
Sequence 15091	AA551358	EST	2321610
Sequence 15092	AA551385	EST	2321637
Sequence 15093	AA551581	EST	2321833

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<u>Table 3-5</u>

Sequence 15094	AA551709	EST	2321961
Sequence 15095	AA551728	EST	2321980
Sequence 15096	AA551770	EST	2322022
Sequence 15097	AA552169	EST	2322421
Sequence 15098	AA552290	EST	2322542
Sequence 15099	AA554025	EST	2324564
Sequence 15100	AA554718	EST	2325257
Sequence 15101	AA554874	EST	2325413
Sequence 15102	AA555194	EST	2325733
Sequence 15103	AA555326	EST	2325865
Sequence 15104	AA557229	EST	2327706
Sequence 15105	AA557760	EST	2328237
Sequence 15106	AA557768	EST	2328245
Sequence 15107	AA563778	EST	2335417
Sequence 15108	AA566072	EST	2337711
Sequence 15109	AA568232	EST	2341286
		EST	2344096
Sequence 15110	AA570116		
Sequence 15111	AA570171	EST	2344151
Sequence 15112	AA570649	EST	2344629
Sequence 15113	AA570705	EST	2344685
Sequence 15114	AA573315	EST	2347843
Sequence 15115	AA573354	EST	2347882
Sequence 15116	AA573559	EST	2348087
Sequence 15117	AA575939	EST	2350454
Sequence 15118	AA576506	EST	2354006
Sequence 15119	AA578782	EST	2356966
Sequence 15120	AA579375	EST	2357559
Sequence 15121	AA579617	EST	2357801
Sequence 15122	AA579690	EST	2357874
Sequence 15123	AA579834	EST	2355161
Sequence 15124	AA580489	EST	2355816
Sequence 15125	AA582370	EST	2359730
Sequence 15126	AA586421	EST	2398416
	AA588541	EST	2401716
Sequence 15127			2401710
Sequence 15128	AA593040	EST	
Sequence 15129	AA595014	EST	2410364
Sequence 15130	AA595339	EST	2410689
Sequence 15131	AA599268	EST	2432893
Sequence 15132	AA599661	LOI	2433286
Sequence 15133	AA600340	EST	2433965
Sequence 15134	AA600781	EST	2434406
Sequence 15135	AA602357	EST	2436335
Sequence 15136	AA604839	EST	2445 <b>70</b> 3
Sequence 15137	AA604894	EST	2445758
Sequence 15138	AA608917	EST	2457345
Sequence 15139	AA608990	EST	2457418
Sequence 15140	AA609219	EST	2457647
Sequence 15141	AA613550	EST	2464588
Sequence 15142	AA613637	EST	2464675
Sequence 15143	AA618493	EST	2505698
Sequence 15144	AA620307	EST	2524246
Sequence 15145	AA620465	EST	2524404
	AA621196		2525135
Sequence 15146		EST	
Sequence 15147	AA622806	EST	2526682
Sequence 15148	AA625155	EST	2537540
Sequence 15149	AA625541	EST	2537928

Page 91 of 105

# <u>Table 3-5</u>

Sequence 15150	AA626377	EST	2538764
Sequence 15151	AA626635	EST	2539022
Sequence 15152	AA626658	EST	2539045
Sequence 15153	AA628413	EST	2540800
Sequence 15154	AA629923	EST	2552534
Sequence 15155	AA629979	EST	2552590
Sequence 15156	AA630335	EST	2552946
Sequence 15157	AA631646	EST	
Sequence 15158	AA631803	EST	2554257
Sequence 15159	AA631901	EST	2554414
Sequence 15160	AA632126	EST	2554512
Sequence 15161	AA632225	EST	2555540
Sequence 15162	AA632270	EST	2555639
Sequence 15163	AA632610		2555684
Sequence 15164	AA633438	EST EST	2556024
Sequence 15165	AA633491	EST	2555298
Sequence 15166	AA634087	EST	2555351
Sequence 15167	AA634119		2557301
Sequence 15168	AA635616	EST EST	2557333
Sequence 15169	AA639362	-	2559458
Sequence 15170	AA640626	EST	2563141
Sequence 15171	AA641037	EST	2565876
Sequence 15172	AA641660	EST	2566287
Sequence 15173	AA644009	EST	2566878
Sequence 15174	AA644184	EST	2569227
Sequence 15175	AA648402	EST EST	2569402
Sequence 15176	AA649327	EST	2574831
Sequence 15177	AA649926		2575756
Sequence 15178	AA649975	EST EST	2577254
Sequence 15179	AA651721	EST	2577303
Sequence 15180	AA652395	EST	2583373 2584047
Sequence 15181	AA659388	EST	2595542
Sequence 15182	AA659693	EST	2595847
Sequence 15183	AA661905	EST	2615996
Sequence 15184	AA662530	EST	2616621
Sequence 15185	AA663778	EST	2617769
Sequence 15186	AA665413	EST	2620026
Sequence 15187	AA669980	EST	2631479
Sequence 15188	AA676967	EST	2657489
Sequence 15189	AA677766	EST	2658288
Sequence 15190	AA679533	EST	2660055
Sequence 15191	AA683488	EST	2670086
Sequence 15192	AA687192	EST	2674397
Sequence 15193	AA687631	EST	2674537
Sequence 15194	AA687667	EST	2674573
Sequence 15195	AA689603	EST	2690530
Sequence 15196	AA699712	EST	2702675
Sequence 15197	AA699807	EST	2702770
Sequence 15198	AA702016	EST	2705129
Seguence 15199	AA702170	EST	2705283
Sequence 15200	AA705025	EST	2714943
Sequence 15201	AA707310	EST	2717228
Sequence 15202	AA708327	EST	2718245
Sequence 15203	AA714016	EST	2726290
Sequence 15204	AA714225	EST	2726499
Sequence 15205	AA716100	EST	2728374

### <u>Table 3-5</u>

Sequence 15206	AA716314	EST	2728588
Sequence 15207	AA720680	EST	2736815
Sequence 15208	AA720790	EST	2736925
Sequence 15209	AA721701	EST	2737763
Sequence 15210	AA723194	EST	2740971
	AA725472		2743179
Sequence 15211		EST	
Sequence 15212	AA725821	EST	2743528
Sequence 15213	AA728892	EST	2750251
Sequence 15214	AA728961	EST	2750320
Sequence 15215	AA729381	EST	2750740
Sequence 15216	AA731128	EST	2752332
Sequence 15217	AA731392,	EST	2753548
Sequence 15218	AA731610	EST	2753766
Sequence 15219	AA733210	EST	2775319
Sequence 15220	AA736880	EST	2768114
Sequence 15221	AA742380	EST	2784380
Sequence 15222	AA742661	EST	2782167
Sequence 15223	AA744728	EST	2783492
Sequence 15224	AA744747	EST	2783511
Sequence 15225	AA744963	EST	2783727
Sequence 15226	AA745648	EST	2785634
Sequence 15227	AA746967	EST	2786925
Sequence 15228	AA747290	EST	2787248
Sequence 15229	AA747733	EST	2787691
Sequence 15230	AA749010	EST	2788968
Sequence 15231	AA749335	EST	2789293
Sequence 15232	AA764788	EST	2816026
Sequence 15232	AA765778	EST	2817016
Sequence 15234	AA768205	EST	2819220
Sequence 15235	AA769138	EST	2820376
Sequence 15236	AA769353	EST	2820591
Sequence 15237	AA769538	EST	2820776
Sequence 15238	AA769986	EST	2821224
Sequence 15239	AA770216	EST	2821454
Sequence 15240	AA770613	EST	2821851
Sequence 15241	AA771700	EST	2823483
	AA771700 AA774100	EST	2825989
Sequence 15242	AA774100 AA775415	EST	2834749
Sequence 15243	AA775624	EST	2834958
Sequence 15244	AA778356	EST	2837687
Sequence 15245			2837991
Sequence 15246	AA778660	EST	2838256
Sequence 15247	AA778925	EST	
Sequence 15248	AA779599	EST	2838930
Sequence 15249	AA780299	EST	2839630
Sequence 15250	AA782060	EST	2841391
Sequence 15251	AA782690	EST	2842021
Sequence 15252	AA805264	EST	2874014
Sequence 15253	AA806729	EST	2875479
Sequence 15254	AA807199	EST	2876775
Sequence 15255	AA807824	EST	2877230
Sequence 15256	AA809127	EST	2878533
Sequence 15257	AA810277	EST	2879636
Sequence 15258	AA812398	EST	2882009
Sequence 15259	AA814392	EST	2883988
Sequence 15260	AA814728	EST	2884324
Sequence 15261	AA814888	EST	2884484

**Table 3-5** 

Sequence 1526	2 AA815422	EST	2885018
Sequence 1526		EST	2896535
Sequence 1526		EST	289 <b>9289</b>
Sequence 1526		EST	2901317
Sequence 1526		EST	
•			2900168
Sequence 1526		EST	2900179
Sequence 1526		EST	2900365
Sequence 1526		EST	2900366
Sequence 1527		EST	2901720
Sequence 1527		EST	2905025
Sequence 1527		EST	2905047
Sequence 1527		EST	2898667
Sequence 1527		EST	29 <b>08787</b>
Sequence 1527		EST	291 <b>0067</b>
Sequence 1527		EST	2910188
Sequence 1527		EST	291 <b>2327</b>
Sequence 1527	8 AA838535	EST	2913334
Sequence 15279		EST	29 <b>297</b> 15
Sequence 1528	0 AA843523	EST	2930041
Sequence 1528	1 AA844956	EST	2931407
Sequence 15282		EST	29 <b>32875</b>
Sequence 1528		EST	29 <b>32896</b>
Sequence 15284	4 AA846815	EST	29 <b>32955</b>
Sequence 1528!	5 AA8539 <b>37</b>	EST	2941475
Sequence 15286	3 AA858162	EST	2946464
Sequence 15287	7 AA861557	EST	295 <b>3697</b>
Sequence 15288	3 AA863230	EST	29557 <b>09</b>
Sequence 15289	AA863232	EST	2955711
Sequence 15290	AA863284	EST	295 <b>5763</b>
Sequence 15291	AA864497	EST	2958810
Sequence 15292	2 AA864573	EST	2958886
Sequence 15293	3 AA865331	EST	2957 <b>607</b>
Sequence 15294	AA8655 <b>52</b>	EST	295 <b>7828</b>
Sequence 15295	5 AA865871	EST	2958147
Sequence 15296	AA866156	EST	2958432
Sequence 15297	AA868706	EST	2964151
Sequence 15298	3 AA8731 <b>73</b>	EST	2969295
Sequence 15299	AA8731 <b>75</b>	EST	2969 <b>297</b>
Sequence 15300	AA877181	EST	298 <b>6258</b>
Sequence 15301	AA877762	EST	2986727
Sequence 15302		EST	2992812
Sequence 15303	AA885293	EST	2994370
Sequence 15304	AA885297	EST	2994374
Sequence 15305	AA886033	EST	3001141
Sequence 15306	AA887219	EST	3002327
Sequence 15307		EST	3017224
Sequence 15308		EST	3017279
Sequence 15309		EST	3017328
Sequence 15310		EST	3017355
Sequence 15311		EST	3038097
Sequence 15312		EST	3040485
Sequence 15313		EST	3041473
Sequence 15314		EST	3047 <b>593</b>
Sequence 15315		EST	3052431
Sequence 15316		EST	3058500
Sequence 15317		EST	3075 <b>842</b>
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### <u>Table 3-5</u>

Sequence 15	374	AI073499	EST	3400143
Sequence 15		AI078597	EST	3413005
Sequence 15		AI081106	EST	3417898
Sequence 15	377	AI081419	EST	3418211
Sequence 15		AI081478	EST	3418270
Sequence 15		AI081536	EST	3418328
Sequence 15		AI084607	EST	3423030
Sequence 15		AI086937	EST	3425360
Sequence 15		AI087840	EST	3426873
Sequence 15		AIC89289	EST	3428348
Sequence 15		AI089629	EST	3428688
Sequence 15		Al091340	EST	3430399
Sequence 15		Al091999	EST	3427197
Sequence 15		AI092232	EST	3431226
		AIC92931	EST	3431907
Sequence 15		AIC93444	EST	3432420
Sequence 15				
Sequence 15		AIC94174 AIC95534	EST EST	3433150
Sequence 15				3434510
Sequence 15		A1095786	EST	3434762
Sequence 15		AI096685	EST	3446179
Sequence 15		Al114501	EST	6359846
Sequence 15		Al114683	EST	6360028
Sequence 15		Al124954	EST	3593468
Sequence 15		Al125449	EST	3593963
Sequence 15		Al125479	EST	3593993
Sequence 15		Al125915	EST	3594429
Sequence 15		Al127049	EST	3595563
Sequence 15		Al127824	EST	3596338
Sequence 15		Al129324	EST	3597838
Sequence 15		Al129703	EST	3598217
Sequence 15		Al129827	EST	3598341
Sequence 15		Al130744	EST	3600760
Sequence 15		Al130885	EST	3600901
Sequence 15		Al132921	EST	6360237
Sequence 15		Al133690	EST	6361006
Sequence 15		Al138441	EST	3644413
Sequence 15		Al139393	EST	3645365
Sequence 15		Al141041	EST	3648498
Sequence 15		Al142953	EST	3659312
Sequence 15		Al143184	EST	3664993
Sequence 15		Al143226	EST	3665035
Sequence 15		Al143638	EST	3665447
Sequence 15		Al143894	EST	3665703
Sequence 15		Al144216	EST	3666025
Sequence 15		Al144385	EST	3666194
Sequence 15		Al146512	EST	3674194
Sequence 15		Al150844	EST	3679313
Sequence 15		Al150878	EST	3679347
Sequence 15		Al159872	EST	3693252
Sequence 15		Al159999	EST	3693379
Sequence 15		Al160331	EST	3693711
Sequence 15		Al160430	EST	3693810
Sequence 15		Al161162	EST	3694467
Sequence 15		Al167145	EST	3700315
Sequence 15		Al167 <b>693</b>	EST	3700863
Sequence 15	429	Al168090	EST	3701260

**Table 3-5** 

Sequence 1543		EST	3721247
Sequence 1543	1 Al174722	EST	6361100
Sequence 1543		EST	6361229
Sequence 1543	3 Al188128	EST	3739337
Sequence 1543	4 Al188187	EST	3739396
Sequence 1543	5 Al188575	EST	3739784
Sequence 1543	6 Al188 <b>768</b>	EST	3739977
Sequence 1543	7 Al189 <b>569</b>	EST	3740778
Sequence 1543	8 Al191342	EST	3742551
Sequence 1543		EST	3743142
Sequence 1544		EST	3751835
Sequence 1544		EST	3754465
Sequence 1544		EST	3755729
Sequence 1544		EST	6361448
Sequence 1544		EST	6361456
Sequence 1544		EST	3770385
Sequence 1544		EST	3801146
Sequence 1544		EST	3804440
Sequence 1544		EST	3804627
Sequence 1544		EST	3804985
Sequence 1545		EST	3805009
Sequence 1545		EST	3836306
Sequence 1545		EST	3836609
Sequence 1545		EST	3837160
Sequence 1545		EST	3839584
Sequence 1545		EST	3841284
Sequence 15456		EST	3841645
Sequence 15457		EST	3844020
Sequence 15458		EST	3844543
Sequence 15459 Sequence 15460		EST EST	3846209
Sequence 1546		EST	3849584 3850451
Sequence 15462		EST	3850369
Sequence 1546		EST	3858537
Sequence 15464		EST	3861885
Sequence 15465		EST	3869780
Sequence 15466		EST	3871576
Sequence 15467		EST	3872208
Sequence 15468		EST	3872492
Sequence 15469		EST	3884566
Sequence 15470		EST	3884904
Sequence 15471		EST	3886508
Sequence 15472	Al267471	EST	3886638
Sequence 15473		EST	3887127
Sequence 15474	Al268542	EST	3887709
Sequence 15475	Al269537	EST	3888704
Sequence 15476	Al271234	EST	3890401
Sequence 15477	Al271608	EST	3890775
Sequence 15478	Al274628	EST	3896896
Sequence 15479	Al276950	EST	3899218
Sequence 15480		EST	3900479
Sequence 15481		EST	3900585
Sequence 15482	Al278411	EST	3900679
Sequence 15483	Al278998	EST	3917232
Sequence 15484		EST	3917734
Sequence 15485	Al279903	EST	3918137

### Table 3-5

Sequence 1	5486	Al281560	EST	3919793
Sequence 1		Al282246	EST	3920479
Sequence 1		Al283100	EST	3921333
Sequence 1		Al284545	EST	3922778
Sequence 1		Al284565	EST	3922798
Sequence 1		Al289108	EST	3932372
Sequence 1	5492	Al289752	EST	3931593
Sequence 1		Al290104	EST	3931770
Sequence 1		Al291500	EST	3934274
Sequence 1		Al291613	EST	3934387
Sequence 1		Al2993B4	EST	3959038
Sequence 1		Al299392	EST	3959046
Sequence 1		Al300346	EST	3959692
		Al300555	EST	3959901
Sequence 1		Al301294	EST	3960640
Sequence 1	5500		EST	3961250
Sequence 1		Al301904		
Sequence 1		Al305197	EST	3988838
Sequence 1		Al305516	EST	3990407
Sequence 1		Al307736	EST	4001954
Sequence 1		Al311614	EST	4006485
Sequence 1		Al313335	EST	4018940
Sequence 1		Al332968	EST	4069527
Sequence 1		Al332994	EST	4069553
Sequence 1		Al334291	EST	4070850
Sequence 1		Al335537	EST	4072464
Sequence 1		Al339426	EST	4076353
Sequence 1		Al339635	EST	4076562
Sequence 1		Al339952	EST	4076879
Sequence 1		Al341076	EST	4078003
Sequence 1		Al342005	EST	4078932
Sequence 1	5516	Al342217	EST	4079144
Sequence 1		Al342486	EST	4079413
Sequence 1		Al344424 ·	EST	4081630
Sequence 1		Al345709	EST	4082915
Sequence 1		Al347461	EST	4084667
Sequence 1		Al348001	EST	4085207
Sequence 1		Al348771	EST	4085989
Sequence 1		Al349772	EST	4086978
Sequence 1	5524	Al352497	EST	4089703
Sequence 1	<b>552</b> 5	Al354464	EST	4094617
Sequence 1	5526	Al356534	EST	4108155
Sequence 1	5527	Al361046	EST	4112667
Sequence 1	5528	Al361513	EST	4113134
Sequence 1	5529	Al362439	EST	4114060
Sequence 1	5530	Al366870	EST	4136615
Sequence 1	5531	Al367932	EST	4137677
Sequence 1	5532	Al368934	EST	4147687
Sequence 1		Al369035	EST	4147788
Sequence 1		Al371024	EST	4149777
Sequence 1		Al371230	EST	4149983
Sequence 1		Al375148	EST	4175138
Sequence 1		Al375903	EST	4175893
Sequence 1		Al376130	EST	4176120
Sequence 1		Al376380	EST	4186229
Sequence 1		Al376921	EST	4186774
Sequence 1		Al377037	EST	4186890
Coquerios 1	JJ71			00000

<u>Table 3-5</u>

Sequence 15598	Al565340	EST	4523797
Sequence 15599	AI566010	EST	4524462
Sequence 15600	AI566802	EST	4525254
Sequence 15601	AI567133	EST	4525585
Sequence 15602	AI567453	EST	4525905
Sequence 15603	AI567945	EST	4526397
Sequence 15604	AI569819	EST	4533193
Sequence 15605	AI569945	EST	4533319
Sequence 15606	AI571329	EST	4534703
Sequence 15607	AI571458	EST	4534832
Sequence 15608	Al581200	EST	4565576
Sequence 15609	Al582529	EST	4568426
Sequence 15610	AI582700	EST	4568597
Sequence 15611	AI597662	EST	4606710
Sequence 15612	Al610362	EST	
Sequence 15613	Al619754	EST	4619529
Sequence 15614	Al620259	EST	4628880 4629385
Sequence 15615	Al621173	EST .	4630299
Sequence 15616	A1624454	EST	
Sequence 15617	Al628617	EST	4649385
Sequence 15618	Ai632017	EST	4665417
Sequence 15619	Al632109	EST	4683347
Sequence 15620	AI632109 AI633086	EST	4683439
Sequence 15621	Al633612		4684416
Sequence 15622	Al634293	EST EST	4684942
Sequence 15623	Al635449	EST	4685623
Sequence 15624	Al636182	EST	4686779
Sequence 15625	AI636818	EST	4687512
Sequence 15626	AI638334	EST	4688148 4690568
Sequence 15627	Al650736	EST	4734715
Sequence 15628	Al651228	EST	4735207
Sequence 15629	Al651423	EST	4735402
Sequence 15630	Al651529	EST	4735508
Sequence 15631	AI652336	EST	4736315
Sequence 15632	Al653693	EST	4737672
Sequence 15633	Al653819	EST	4737798
Sequence 15634	A1654736	EST	4738715
Sequence 15635	AI655590	EST	4739569
Sequence 15636	AI655840	EST	4739819
Sequence 15637	AI659386	EST	4762956
Sequence 15638	AI660901	EST	4764484
Sequence 15639	Al671089	EST	4850820
Sequence 15640	AI673189	EST	4852920
Sequence 15641	Al674314	EST	4874794
Sequence 15642	Al674456	EST	4874936
Sequence 15643	Al674918	EST	4875398
Sequence 15644	AI677810	EST	4887992
Sequence 15645	AI677876	EST	4888058
Sequence 15646	AI679799	EST	4889981
Sequence 15647	A1680555	EST	4890737
Sequence 15648	AI680742	EST	4890924
Sequence 15649	AI681316	EST	4891498
Sequence 15650	A1683072	EST	4893254
Sequence 15651	AI683117	EST	4893299
Sequence 15652	AI683727	EST	4893909
Sequence 15653	AI689427	EST	4900721

**Table 3-5** 

Sequence 15710	AI961563	EST	5754276
Sequence 15711	AI9 <b>71397</b>	EST	5768223
Sequence 15712	AI98 <b>6177</b>	EST	5813454
Sequence 15713	AL036349	EST	<b>5</b> 4059 <b>30</b>
Sequence 15714	AL03 <b>7069</b>	EST	<b>59</b> 280 <b>30</b>
Sequence 15715	AL03 <b>7845</b>	EST	5407176
Sequence 15716	AL037978	EST	5407290
Sequence 15717	AL038564	EST	5407750
Sequence 15718	AL039253	EST	5408327
Sequence 15719	AL040602	EST	<b>5</b> 40954 <b>9</b>
Sequence 15720	AL042384	EST	5421723
Sequence 15721	AL046811	EST	5434873
Sequence 15722	AL04 <b>9135</b>	EST	4728445
Sequence 15723	AL118571	EST	5924470
Sequence 15724	AL120354	EST	<b>59</b> 2625 <b>3</b>
Sequence 15725	AL133982	EST	6602169
Sequence 15726	AL134985	EST	6603172
Sequence 15727	AL138451	EST	6855132
Sequence 15728	AL157600	EST	7058001
Sequence 15729	AW008029	EST	<b>58</b> 56807
Sequence 15730	AW00 <b>8673</b>	EST	5857451
Sequence 15731	AW009152	EST	5857930
Sequence 15732	AW016461	EST	<b>586</b> 52 <b>18</b>
Sequence 15733	AW020561	EST	5874091
Sequence 15734	AW020914	EST	5874444
Sequence 15735	AW021803	EST	<b>587</b> 53 <b>33</b>
Sequence 15736	AW022479	EST	<b>587</b> 60 <b>09</b>
Sequence 15737	AW023 <b>580</b>	EST	5877110
Sequence 15738	AW02 <b>6439</b>	EST	<b>587</b> 99 <b>69</b>
Sequence 15739	AW026 <b>700</b>	EST	5880153
Sequence 15740	AW027717	EST	5886473
Sequence 15741	AW029611	EST	5888367
Sequence 15742	AW051380	EST	<b>5</b> 913650
Sequence 15743	AW055331	EST	5921034
Sequence 15744	AW057525	EST	5933164 '
Sequence 15745	AW062700	EST	6014085
Sequence 15746	AW069166	EST	6024252
Sequence 15747	AW075351	EST	6030349
Sequence 15748	AW079672	EST	6034824
Sequence 15749	AW081110	EST	6036262
Sequence 15750	AW081275	EST	6036427
Sequence 15751	AW083877	EST	6039029
Sequence 15752	AW084353	EST	6039505
Sequence 15753	AW084441	EST	6039593
Sequence 15754	AW085581	EST	6040733
Sequence 15755	AW088099	EST	6043904
Sequence 15756	AW089689	EST	6047033
Sequence 15757	AW 102924	EST	6073539
Sequence 15758	AW 131444	EST	6133051
Sequence 15759	AW 151003	EST	6198901
Sequence 15760	AW151170	EST	6199068
Sequence 15761	AW161328	EST	6300361
Sequence 15762	AW161668	EST	6300701
Sequence 15763	AW166819	EST	<b>63</b> 98344
Sequence 15764	AW167129	EST	6398654
Sequence 15765	AW 168050	EST	<b>639</b> 95 <b>75</b>

<u>Table 3-5</u>

Sequence 15766	AW177001	EST	6443038
Sequence 15767	AW183074	EST	645 <b>1536</b>
Sequence 15768	AW188081	EST	6462517
Sequence 15769	AW188641	EST	6463001
Sequence 15770	AW188668	' EST	6463028
Sequence 15771	AW188775	EST	64 <b>63211</b>
Sequence 15772	AW188814	EST	64 <b>63250</b>
Sequence 15773	AW190568	EST	6465048
Sequence 15774	AW235964	EST	65 <b>68353</b>
Sequence 15775	AW243325	EST	65 <b>77165</b>
Sequence 15776	AW243940	EST	6577780
Sequence 15777	AW271299	EST	6658329
Sequence 15778	AW271852	EST	6658882
Sequence 15779	AW273750	EST	6660780
Sequence 15780	AW290969	EST	6697605
Sequence 15781	AW293305	EST	6699867
Sequence 15782	AW293616	EST	6700 <b>252</b>
Sequence 15783	AW294158	EST	6700794
Sequence 15784	AW297885	EST	6704510
Sequence 15785	AW328141	EST	67 <b>98637</b>
Sequence 15786	AW360820	EST	68 <b>65470</b>
Sequence 15787	AW361122	EST	6865772
Sequence 15788	AW365134	EST	6869784
Sequence 15789	AW368376	EST	6873026
Sequence 15790	AW369111	EST	6873761
Sequence 15791	AW378964	EST	6883623
Sequence 15792	AW381285	EST	6885944
Sequence 15793	AW386920	EST	6891579
Sequence 15794	AW390476	EST	6895135
Sequence 15795	AW394098	EST	6898757
Sequence 15796	AW 402596	EST	6921298
Sequence 15797	AW449808	EST	6990584
Sequence 15798	AW449890	EST	6990666
Sequence 15799	AW450105	EST	6990881
Sequence 15800	AW450574	EST	6991350
Sequence 15801	AW450748	EST	6991524
Sequence 15802	AW451232	EST	6992008
Sequence 15803	AW451555	EST	69 <b>92331</b>
Sequence 15804	AW451783	EST	69 <b>92559</b>
Sequence 15805	AW452088	EST	6992864
Sequence 15806	AW452730	EST	6993506
Sequence 15807	AW468168	EST	7038274
Sequence 15808	AW469281	EST	703 <b>9387</b>
Sequence 15809	AW470264	EST	7040370
Sequence 15810	AW473121	EST	7043227
Sequence 15811	AW510526	EST	N/A
Sequence 15812	AW511443	EST	N/A
Sequence 15813	AW572897	EST	7237630
Sequence 15814	C00019	EST	1432249
Sequence 15815	C14194	EST	1568901
Sequence 15816	. C16523	EST	1571230
Sequence 15817	D53484	EST	95 <b>5381</b>
Sequence 15818	D61456	EST	97 <b>0232</b>
Sequence 15819	D61516	EST	970472
Sequence 15820	D81678	EST	1179555
Sequence 15821	F08667	EST	672530
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Page 103 of 105

Table 3-5

Sequence 15822	F10548	EST	683206
Sequence 15823	F35428	EST	4821054
Sequence 15824	H19093	EST	885333
Sequence 15825	H22094	EST	890789
Sequence 15826	H22238	EST	890933
	H39906	EST	915958
Sequence 15827			
Sequence 15828	H47418	EST	923470
Sequence 15829	H52272	EST	992113
Sequence 15830	H52374	EST	992215
Sequence 15831	H54240	EST	994387
Sequence 15832	H64134	EST	1018935
Sequence 15833	H67372	EST	1026112
Sequence 15834	H67494	EST	1026234
Sequence 15835	H77381	EST	1055470
Sequence 15836	H78888	EST	1056977
Sequence 15837	H79965	EST	1058054
Sequence 15838	H81340	EST	1059429
Sequence 15839	H90127	EST	1080557
Sequence 15840	N25954	EST	1140302
Sequence 15841	N43012	EST	1166756
Sequence 15842	N48918	EST	1190084
Sequence 15843	N56877	EST	1200767
Sequence 15844	N62671	EST	1210500
Sequence 15845	N66909	EST	1219034
Sequence 15846	N71827	EST	1228539
Sequence 15847	N95674	EST	1267942
Sequence 15848	N95803	EST	1268098
Sequence 15849	R23565	EST	778453
Sequence 15850	R80268	EST	856549
Sequence 15851	R93 <b>800</b>	EST	967966
Sequence 15852	R98534	EST	985051
Sequence 15853	T49212	EST	651072
Sequence 15854	T55511	EST	657372
Sequence 15855	T75284	EST	692046
Sequence 15856	T81310	EST	704195
Sequence 15857	T84643	EST	712995
Sequence 15858	T91068	EST	722981
Sequence 15859	U69567	EST	2731398
Sequence 15860	W02490	EST	1274488
Sequence 15861	W04263	EST	1276162
Sequence 15862	W05733	EST	1278455
Sequence 15863	W19899	EST	1295768
Sequence 15864	W21207	EST	1298239
Sequence 15865	W25889	EST	1306147
Sequence 15866	W58768	EST	1364280
Sequence 15867	W60245	EST	1367175
Sequence 15868	W61046	EST	1367806
Sequence 15869	W79674	EST	1390063
Sequence 15870	Z22018	EST	288765
Sequence 15871	Z40994	EST	566739
Sequence 15872	T00880	NUCPATENT	N/A
Sequence 15873	T19326	NUCPATENT	N/A
Sequence 15874	T27758	NUCPATENT	N/A
Sequence 15875	T48099	NUCPATENT	N/A
Sequence 15876	T92160	NUCPATENT	N/A
Sequence 15877	T96835	NUCPATENT	N/A
23quanos 13077	. 00000	MOO! WIEM!	17/0

Page 104 of 105

Table 3-5

Sequence 15878	V58586	NUCPATENT	N/A
Sequence 15879	V59067	NUCPATENT	N/A
Sequence 15880	V59630	NUCPATENT	N/A
Sequence 15881	V60015	NUCPATENT	N/A
Sequence 15882	V65326	NUCPATENT	N/A
Sequence 15883	<b>V88</b> 598	NUCPATENT	N/A
Sequence 15884	V90354	NUCPATENT	N/A
Sequence 15885	X02974	NUCPATENT	N/A
Sequence 15886	X20483	NUCPATENT	N/A
Sequence 15887	X26850	NUCPATENT	N/A
Sequence 15888	X27348	NUCPATENT	N/A
Sequence 15889	X30410	NUCPATENT	N/A
Sequence 15890	X33813	NUCPATENT	N/A
Sequence 15891	X84945	NUCPATENT	N/A
Sequence 15892	X86772	NUCPATENT	N/A
Sequence 15893	X87396	NUCPATENT	N/A
Sequence 15894	X87868	NUCPATENT	N/A
Sequence 15895	X98006	NUCPATENT	N/A
Sequence 15896	X98010	NUCPATENT	N/A
Sequence 15897	X98409	NUCPATENT	N/A
Sequence 15898	X98714	NUCPATENT	N/A
Sequence 15899	Z14927	NUCPATENT	N/A
Sequence 15900	Z16190	NUCPATENT	N/A
Sequence 15901	Z18356	NUCPATENT	N/A
Sequence 15902	Z24811	NUCPATENT	N/A
Sequence 15903	Z29718	NUCPATENT	N/A
Sequence 15904	Z33656	NUCPATENT	N/A
Sequence 15905	Z34089	NUCPATENT	N/A
Sequence 15906	Z41384	NUCPATENT	N/A
Sequence 15907	<b>Z</b> 97274	NUCPATENT	N/A
Sequence 15908	AC28082	PREPATNUC	N/A
Sequence 15909	AC34864	PREPATNUC	N/A
Sequence 15910	AC36253	PREPATNUC	N/A
Sequence 15911	AC3B032	PREPATNUC	N/A

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANN CCNGGCCCNCCCNCTGGNGNNTTCCCANATCCCNNTTTNTGAATTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** Sequence 11677: Found in patent publication WO99/46375 AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATITAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206 AAAAAAAAAAAAAAAAAAAAAAAA Sequence 11679: Found in patent publication WO99/63088 TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT GACCAGTA Sequence 11680: Found in patent publication WO99/67383 AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

Page 1 of 2

CATCTCAAAAATAAATAAATATATAAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTITGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT / GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAATAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTCTTTTTAGACTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AACAACAAAAAAAACNAAAAATANAAAAAAAAAAAAAANAANNANNCAANNNAAAANTAA AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGATGAGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC
TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT
TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC
TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

Page 1 of 2

CATCTCAAAAATAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATT.TTCCTT TCTCACATAATTACTTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AACAACAAAAAAACNAAAAATANAAAAAAAAAAAAANAANNNANNCAANNNAAAANTAA AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328  $oldsymbol{A}ol$ Sequence 15910: Found in patent publication WO99/66041  $oldsymbol{A}ol$ Sequence 15911: Found in patent publication WO00/05382 AAGACATT CCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGATGAGGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

Page 2 of 2

Sequence 11679: Found in patent publication WO99/63088

Sequence 11678: Found in patent publication WO98/54206

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC
TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT
TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC
TATATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAAATGTTTAGAA

Sequence 13326; Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

Page 1 of 2

TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AACAACAAAAAAACNAAAAATANAAAAAAAAAAAAAANAANNNANNCAANNNAAAANTAA AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGATGAGGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC
TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT
TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC
TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

Page 1 of 2

CATCTCAAAAATAAATAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AACAACAAAAAAACNAAAAATANAAAAAAAAAAAAANAANNNANNCAANNNAAAANTAA **AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA AAATAANTAAAAAAATTNA**TAAAN**AATAAATNTTAAA**TAAAAAAATATT Sequence 15909: Found in patent publication WO99/14328 **AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA** Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA 

Page 2 of 2

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGCGCCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC NCTCCAGCCTGGGCAACAGAGTGAGACTCCGTCCCGCTAANAAATNANAAAANAANAANA CCNGGCCCNCCCNCTGGNGNNTTCCCANATCCCNNTTTTNTGAATTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TTGTTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** Sequence 11677: Found in patent publication WO99/46375

AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT TTCTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

Sequence 11679: Found in patent publication WO99/63088

AGTACTTCTAGAATTAATTAAGGCGGGGGAGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT **GACCAGTA** 

Sequence 11680: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC ATGCTT

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA 

Sequence 13324: Found in patent publication WO99/66041

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

Page 1 of 2

CATCTCAAAAATAAATATATATAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAATAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

PCT/US01/02318 WO 01/053836

#### **Table 3-6**

Sequence 11676: Found in patent publication WO00/05367 AGTACTTCTAGAATTAATTAAGGCGGGCGCCTGCANTCCCAGNTACNCGGNAGGCTGAGG CAGGAGAATGGCGTGAACCCANGAGGCGGAGCTTGCAGTGAGCTGAGATCGCGCCACTGC CCNGGCCCNCCCNCTGGNGNNTTCCCANATCCCNNTTTTNTGAATTTTTTNCCCCCCCC NNNTGGNCCTGTTTTCTACTAGTGATGATCTGGTAATATACAATTTGTCCAGTAGCCAGT TIGTTTTATTGTGTTTTCTAACCATAAGAGATCATTAAAGGCAAAGCCTGTATGACGCT **AATCTTCTGGTCA** Sequence 11677: Found in patent publication WO99/46375

AGTACTTCTAGAATTAATTAAGGCGGGGACACCACTTTTCAAAGGACTTCTTGGTTTCAG CATAACCTAAGACAGGGAATTGGGAGCCATCATATGTCACAGTGTTCAGAATTCAAGCAT TACATTTAGCTGACAAAGTGCTTATAGAATTTCTTAACAACTGAACCATTCAAAAGGATT TCCATAGCTTCTTATCCCTGTATTGAGGTAATAAATTGTTTTACTGACAATTTTTCCTTT

TICTACACTAAAACAATATGTGATATATTTCCCCTCTTGAAGAGGCAATTCATTAAACTC Sequence 11678: Found in patent publication WO98/54206

AAAAAAAAAAAAAAAAAAAAAAAA

Sequence 11679: Found in patent publication WO99/63088 AGTACTTCTAGAATTAATTAAGGCGGGGAGTTTTATCTATTCTTTACATTGGATGCAAAA TGTATTACTCAAGAAGAGGCATTCGGTATCGAACCATAGATGAACATGATGCCATCATTT GTTTATTAATAGTTTAAAACAATATTCTCTTTTTGAAAATAGTATAAACAGGCCATGCAT ATAATGTACAGTGTATTACGTAAATATGTAAAGATTCTTCAAGGTAACAAGGGTTTGGGT TTTGAAATAAACATCTGGATCTTATAGACCGTTCATACAATGGTTTTAGCAAGTTCATAG TAAGACAAACAAGTCCTATCTTTTTTTTTTTGGCTGGGGTGGGGGCATTGGTCACATAT

Sequence 11680: Found in patent publication WO99/67383
AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC **ATGCTT** 

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA 

Sequence 13324: Found in patent publication WO99/66041

ΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑΑ

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

Page 1 of 2

CATCTCAAAAATAAATAAATATATATAAAAAAAAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 **AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT** GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG TTCGCACCTGTAGTCCCAGCTACTGGGGGGGGGGCTGAGGCAGGAGAATGGCTTGAACCCGGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT. TCTCACATAATTACTTTTTCTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT CCAATATGAAGAACTGAGCCTGGAGAGATGGGGATGAGGGGGGAGTGTCGAACCTTTTAGGC TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGGCTCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

Sequence 11677: Found in patent publication WO99/46375

Sequence 11679: Found in patent publication WO99/63088

Sequence 11680: Found in patent publication WO99/67383

Sequence 13322: Found in patent publication WO00/09552

Sequence 13323: Found in patent publication WO99/66041

AGTACTTCTAGAATTAATTAAGGCGGGGGAAAGTTAGAGGAACTGAAAGTTTGGGAATAG GCTGACCACATATTATGCCAGTGACCAGTATGACAGGAGATGGGGCCCTGCTGCCAGTCA

Sequence 13325: Found in patent publication WO99/67383

AGTACTTCTAGAATTAATTAAGGCGGGTCCCTGCTTGTCCTATGAACTGCTCAGAGCTCC TGTCAGTCCAGCTGGGCCTTCTGGGTTCTGGCACCATTTCGTAGCCATTCTCTTTGTATT TTAAAAGGACGTTATGAAAGGGCTTAGACCAAAATAAATCATAATGTTACTTGAGCCACC TTATATAGCTGCTTGGAGAGTCTGTGTAGTTCTTTCTGCATGTATTAAAAAATGTTTAGAA

Sequence 13326: Found in patent publication WO00/06698

GCCCTGGGAAACGATTGTGAACGCNTGAATGAATTGATGACTAANATCCGCTGCGGGGGT CCTACAGCGCANATGGTAATGCCCGTTCTGACTGGCTGGGAACGGCACCTTAGCAAGATA CTTAAAAGGCGCCTTCTGTGTGCCACNTGCACTCCACCCTGGGCAACAGAGCAAANACCC

CATCTCAAAAATAAATAAATATATAAAAAAAATAAAAAGCTATTTCTAGTTTNATTTCAC TATAAAGTTTTGCTTTTATTAAA Sequence 13327: Found in patent publication US6034229 AGTACTTCTAGAATTAATTAAGGCGGGCGGATCATGAGGTCAGGAGTTCGAGACCAGCCT GGCCAACATGGTGAAACTCCGTGTCTACTAAAAATACAAAAATTGGCTGGGCCGTGGTGG AGGGTAGAGGTTGCAGTGAGCCAAGATTGTGCCACTGCACTCCAGCCTGGGCGACTGAGC AAGACTGTCTAAAAAAAAAAAAAAAATAAATAAAGNAA Sequence 13328: Found in patent publication WO00/18925 GCCGTCGTTTTACACCCCGNAANAANCNATANCNATGATNGNTGCCNGTCCCNCTCNNNA NAGNAATAAAAANTAATTAANGCGGGTGATTAAATTTCAGTCCATTATGATTTTCCTT TCTCACATAATTACTTTTTTTTTTTTTTTTTAGACTTATAAGCTAGCAATTACAGATTTAACTAC **AGCTATCAGCATGGACAAAAATAGTTATACAGCATTTTATAACAGAGCATTATGTTACAC** CAAGATAAGGGAACTTCAAATGGCATTAACAGATTATGGAATTGTGCTGCTTCTTGATGC TACAGAAACTGTNAAACTAAATACCTTCCTTAATCGTGGACTCATCTACGTAGAACTAGG CCAGTATGGCTTTGCACTAGAGGATTTTAAAAC Sequence 15908: Found in patent publication WO98/33916 AAATTNNNTATTTATTTTAAANTAAAAANTATACTNAACTAATTATTTNAATTAAAGAAA Sequence 15909: Found in patent publication WO99/14328 Sequence 15910: Found in patent publication WO99/66041 Sequence 15911: Found in patent publication WO00/05382 AAGACATTCCAGACAAAAAGACTAACTTGTCAGAAAGCCCTGTGGCGGAAGGGAGCTTTT TTTGTAAAGGAGTTTTGGTTTTCTCCTAATAGCAATGGGATATCTTCCAAGGAATCTCAA TCAAAAGGGAGAGATGCCCGATTGGAATGTCATCCCTGGCTGAAGAGTAGAGGAAGCA AAAAAAAAAAAAAAAAAAAAANNGNNNNNNCCCNNTNTTTTTTT

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Sequence 7	AL039430	Sequence 59	Al267285
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Page 1 of 74

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Table 4-1

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Page 2 of 74

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Page 3 of 74

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Page 4 of 74

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Page 5 of 74

Table 4-1

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Page 6 of 74

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Page 7 of 74

Table 4-1

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Sequence 753	AA744849	Sequence 805	AA947511
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Sequence 756	AA228940	Sequence 808	AA324252
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Page 8 of 74

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Sequence 882	AA315621	Sequ <b>ence 934</b>	AA040135
Sequence 883	AA075012	Sequence 935	AA480124

Page 9 of 74

Sequence 936	AA233943	Sequence 988	AI880872
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Page 10 of 74

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Page 11 of 74

Sequence 1144	AA96 <b>8556</b>	Sequence 1196	Z25169
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Page 12 of 74

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Page 13 of 74

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Sequence 1354	AB011158	Sequence 1406	AF141327
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Page 14 of 74

Table 4-1

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Sequence 1478	AF027515	Sequence 1530.	AB023224
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Page 15 of 74

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Sequence 1611	AB018281	Sequence 1663	AF055994

Page 16 of 74

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Sequence 1713	AB018296	Sequence 1765	AF100755
Sequence 1714	M22918	Sequence 1766	AF078860
Sequence 1715	L05186	Sequence 1767	U03886

Page 17 of 74

Sequence 1768	L07033	Sequence 1820	D38555
Sequence 1769	U07919	Sequence 1821	D28540
Sequence 1770	D13629	Sequence 1822	AF191298
Sequence 1771	AF191018	Sequence 1823	AF042346
Sequence 1772	X04697	Sequence 1824	E01956
Sequence 1773	X81109	Sequence 1825	X70394
Sequence 1774	AL079298	Sequence 1826	U41060
Sequence 1775	X56958	Sequence 1827	AB000887
Sequence 1776	AL110126	Sequence 1828	U25182
Sequence 1777	X63679	Sequence 1829	X87241
Sequence 1778	X17620	Sequence 1830	AB007972
Sequence 1779	A21185	Sequence 1831	M63573
Sequence 1780	AF087942	Sequence 1832	X07270
Sequence 1781	E01954 ·	Sequence 1833	AF124439
Sequence 1782	AF077200	Sequence 1834	D50408
Sequence 1783	D87078	Sequence 1835	J04443
Sequence 1784	AL050272	Sequence 1836	D21260
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Sequence 1793	D87666	Sequence 1845	M26481
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Sequence 1796	X02875	Sequence 1848	D14812
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Sequence 1804	AF088004	Sequence 1856	X60221
Sequence 1805	AF086431	Sequence 1857	AB007882
Sequence 1806	AF010233	Sequence 1858	AF082513
Sequence 1807	AF086336	Sequence 1859	AF071202
Sequence 1808	AF057705	Sequence 1860	M26383
Sequence 1809	AB018280	Sequence 1861	X57351
Sequence 1810	AF016496	Sequence 1862	AF007142
Sequence 1811	M26663	Sequence 1863	AF020202
Sequence 1812	M17886	Sequence 1864	AB020686
Sequence 1813	X07466	Sequence 1865	X91257
Sequence 1814	D31767	Sequence 1866	D32051
Sequence 1815	L05779	Sequence 1867	AF147339
Sequence 1816	U07809	Sequence 1868	AB002306
Sequence 1817	AF137334	Sequence 1869	AF100756
Sequence 1818	AF056322	Sequence 1870	U01925
Sequence 1819	AB001636	Sequence 1871	D14878

Page 18 of 74

Sequence 1872	M22590	Sequence 1924	X27262
Sequence 1873	M93651	Sequence 1925	T59274
Sequence 1874	AF054838	Sequence 1926	V69358
Sequence 1875	M69066	Sequence 1927	X98998
Sequence 1876	AF017269	Sequence 1928	Z10658
Sequence 1877	J03007	Sequence 1929	X76578
Sequence 1878	AF019214	Sequence 1930	Z06652
Sequence 1879	AF039023	Sequence 1931	T37414
Sequence 1880	X66276	Sequence 1932	V56670
Sequence 1881	AL110179	Sequence 1933	V58682
Sequence 1882	D86984	Sequence 1934	Z15583
Sequence 1883	D90226	Sequence 1935	T18813
Sequence 1884	AL035304	Sequence 1936	X22116
Sequence 1885	D13643	Sequence 1937	T61701
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Sequence 1887	L28997	Sequence 1939	V58756
Sequence 1888	AB023173	Sequence 1940	V57903
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Sequence 1890	AB033899	Sequence 1942	Z17335
Sequence 1891	U90550	Sequence 1943	X26850
Sequence 1892	M10119	Sequence 1944	Z33658
Sequence 1893	AL133078	Sequence 1945	Z14392
Sequence 1894	J04607	Sequence 1946	V63172
Sequence 1895	J02642	Sequence 1947	Z15729
Sequence 1896	A06919	Sequence 1948	V34236
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Sequence 1900	AF083441	Sequence 1952	AC33865
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Sequence 1903	U02680	Sequence 1955	AC36206
Sequence 1904	L08599	Sequence 1956	AC36220
Sequence 1905	AF019226	Sequence 1957	AC36078
Sequence 1906	AF176574	Sequence 1958	AC36205
Sequence 1907	AB028977	Sequence 1959	AC35218
Sequence 1908	AJ000881	Sequence 1960	AC36180
Sequence 1909	D89678	Sequence 1961	AC36187
Sequence 1910	D50405	Sequence 1962	AC36079
Sequence 1911	D29641	Sequence 1963	AC36823
Sequence 1912	M34840	Sequence 1964	AC36709
Sequence 1913	V62427	Sequence 1965	AA304651
Sequence 1914	V62430	Sequence 1966	AA308091
Sequence 1915	X20432	Sequence 1967	AA442287
Sequence 1916	V87930	Sequence 1968	AA460440
Sequence 1917	V58694	Sequence 1969	AA513597
Sequence 1918	X98608	Sequence 1970	AA576425
Sequence 1919	Z15132	Sequence 1971	AB032983
Sequence 1920	V58628	Sequence 1972	AF047020
Sequence 1921	Q97526	Sequence 1973	AF103907
Sequence 1922	X25445	Sequence 1974	A1080439
Sequence 1923	X04318	Sequence 1975	AI088633

Page 19 of 74

Table 4-1

Sequence 1976	Al267321	Sequence 2028	AA663192
Sequence 1977	Al274700	Sequence 2029	AA449889
Sequence 1978	AI557336	Sequence 2030	AA209369
Sequence 1979	Al825988	, Sequence 2031	AA354730
Sequence 1980	AL037105	Sequence 2032	AA354196
Sequence 1981	AW138181	Sequence 2033	A1791370
Sequence 1982	D14043	Sequence 2034	AA447433
Sequence 1983	L38961	Sequence 2035	N44743
Sequence 1984	M26880	Sequence 2036	AA528104
Sequence 1985	X77548	Sequence 2037	AA853882
Sequence 1986	AB001636	Sequence 2038	AA134726
Sequence 1987	AB008681	Sequence 2039	AA811333
Sequence 1988	AB020866	Sequence 2040	AL121340
Sequence 1989	AC000399	Sequence 2041	AA310201
Sequence 1990	AC002086	Sequence 2042	A1694572
Sequence 1991	AC003991	Sequence 2043	Al140921
Sequence 1992	AC004507	Sequence 2044	AA149579
Sequence 1993	AC004816	Sequence 2045	AA506729
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Sequence 1995	AC005480	Sequence 2047	Al199094
Sequence 1996	AC005538	Sequence 2048	AA975275
Sequence 1997	AC005831	Sequence 2049	N25122
Sequence 1998	AC006153	Sequence 2050	Al636911
Sequence 1999	AL031777	Sequence 2051	AI219803
Sequence 2000	AL117340	Sequence 2052	AA483044
Sequence 2001	AL121603	Sequence 2053	AA522552
Sequence 2002	AL121654	Sequence 2054	AI564822
Sequence 2003	AL122007	Sequence 2055	R63251
Sequence 2004	AL135879	Sequence 2056	AA526369
Sequ <b>enc</b> e 2005	AP000303	Sequence 2057	AI074652
Sequence 2006	M11548	Sequence 2058	AA603861
Sequence 2007	S65371	Sequence 2059	AA253339
Sequence 2008	U13800	Sequence 2060	AA313170
Sequence 2009	U91321	Sequence 2061	AA228940
Sequence 2010	Z63584	Sequence 2062	AF001542
Sequence 2011	Z73900	Sequence 2063	A1090642
Sequence 2012	AA079188	Sequence 2064	AA035559
Sequence 2013	Al267838	Sequence 2065	AA304845
Sequence 2014	AA507791	Sequence 2066	AI652112
Sequence 2015	AA640474	Sequence 2067	AA058381
Sequence 2016	AI023483	Sequence 2068	AA935526
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Sequence 2018	AA325789	Sequence 2070	Al034131
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Sequence 2021	H70269	Sequence 2073	AI267285
Sequence 2022	AA226321	Sequence 2074	AA531386
Sequence 2023	W58683	Sequence 2075	AA493331
Sequence 2024	AL043205	Sequence 2076	AL043046
Sequence 2025	AA056482	Sequence 2077	Al373120
Sequence 2026	H59731	Sequence 2078	AI017239
Sequence 2027	R69940	Sequence 2079	AA149337
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Page 20 of 74

Sequence 2080	AA622482	Sequence 2132	AI269205
Sequence 2081	AA232147	Sequence 2133	AI754013
Sequence 2082	Al453181	Sequence 2134	AA336412
Sequence 2083	AL121262	Sequence 2135	AW148750
Sequence 2084	AA306322	Sequence 2136	AA304846
Sequence 2085	Al635209	Sequence 2137	AA190998
Sequence 2086	AA973620	Sequence 2138	AA564505
Sequence 2087	AA323181	Sequence 2139	A1086059
Sequence 2088	AA399177	Sequence 2140	AA235224
Sequence 2089	R33609	Sequence 2141	AA405663
Sequence 2090	AW149706	Sequence 2142	C21370
Sequence 2091	AA878395	Sequence 2143	AI253335
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Sequence 2093	AA099399	Sequence 2145	AA314893
Sequence 2094	AA467869	Sequence 2146	Al630616
Sequence 2095	AA747909	Sequence 2147	W27229
Sequence 2096	AA483008	Sequence 2148	AA195525
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Sequence 2100	AL046322	Sequence 2152	AA843845
Sequence 2101	AA146981	Sequence 2153	Al204916
Sequence 2102	AA152337	Sequence 2154	AA172043
Sequence 2103	AA280917	Sequence 2155	AI557235
Sequence 2104	AL043257	Sequence 2156	AA020899
Sequence 2105	AA026149	Sequence 2157	Al954167
Sequence 2106	AA732538	Sequence 2158	AI380932
Sequence 2107	AA876439	Sequence 2159	AI267502
Seguence 2108	AA405535	Sequence 2160	H30141
Sequence 2109	H80936	Sequence 2161	AA634408
Sequence 2110	A1378290	Sequence 2162	AA031287
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Sequence 2113	AA703396	Sequence 2165	AA088714
Sequence 2114	AA312076	Sequence 2166	AA523354
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Sequence 2119	AA506459	Sequence 2171	AA716414
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Sequence 2131	AA764981	Sequence 2183	AA703816

Page 21 of 74

Sequence 2184	AA487754		AA531227
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Sequence 2186	AI671394		AA132904
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Sequence 2218	AA857069	Sequence 2270	AA128591
Sequence 2219	AA765712	Sequence 2271	AW028937
Sequence 2220	AA280256	Sequence 2272	AA503943
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Sequence 2230	AA040135	Sequence 2282	AA994993
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Sequence 2234		Sequence 2286	AA442517
Sequence 2235	AA351504	Sequence 2287	AL044723

Page 22 of 74

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Sequence 2338	A1253379	Sequence 2389	AA658393
Sequence 2339	AA584428	Sequence 2390	Al199681 ·
ocquerios 2009	Al147850	Sequence 2391	H20388

Page 23 of 74

Sequence 2392         AI375141         Sequence 2444         AI970919           Sequence 2394         A193086         Sequence 2445         AL039208           Sequence 2395         AA180137         Sequence 2447         AA459455           Sequence 2397         AA101742         Sequence 2448         AA476360           Sequence 2398         AI028097         Sequence 2450         AI028262           Sequence 2400         AA071199         Sequence 2451         H03785           Sequence 2401         AA479862         Sequence 2453         AA305627           Sequence 2401         AA479862         Sequence 2455         AA082342           Sequence 2401         AA31865         Sequence 2453         AA082342           Sequence 2401         AA31865         Sequence 2453         AA086128           Sequence 2401         AA3186757         Sequence 2455         AA086128           Sequence 2405         AA904265         Sequence 2455         AA2280852           Sequence 2407         AA23143         Sequence 2459         AA234740           Sequence 2408         AI093458         Sequence 2459         AA225294           Sequence 2410         AA736613         Sequence 2462         AA226066           Sequence 2410         A				
Sequence 2394 AA183086 Sequence 2446 AI557336 Sequence 2395 AA180137 Sequence 2447 AA459455 Sequence 2396 AA58673 Sequence 2447 AA459455 Sequence 2397 AA101742 Sequence 2449 AI080272 Sequence 2398 AI028097 Sequence 2450 AI025262 Sequence 2399 A865185 Sequence 2451 AI025262 Sequence 2400 AA071199 Sequence 2451 AA305627 Sequence 2400 AA071199 Sequence 2452 AA305627 Sequence 2401 AA479962 Sequence 2453 AA305627 Sequence 2403 AA318652 Sequence 2404 AA1867557 Sequence 2404 AA1867567 Sequence 2405 AA306128 Sequence 2406 AA167677 Sequence 2455 AA086128 Sequence 2406 AL040264 Sequence 2456 AI0823362 Sequence 2407 AA243143 Sequence 2457 AA280852 Sequence 2408 AI093458 Sequence 2409 AA045381 Sequence 2401 AA736613 Sequence 2401 AA736613 Sequence 2402 Sequence 2410 AA736613 Sequence 2462 AA287720 Sequence 2411 W04240 Sequence 2462 AA287720 Sequence 2414 AA748064 Sequence 2415 Sequence 2416 AA736613 Sequence 2465 AA368573 Sequence 2416 AA748064 Sequence 2416 AA736861 Sequence 2417 AA004037 Sequence 2418 AA040437 Sequence 2418 AA040437 Sequence 2418 AA040437 Sequence 2418 AA040437 Sequence 2419 AA366876 Sequence 2419 AA366876 Sequence 2419 AA366876 Sequence 2410 Sequence 2410 Sequence 2411 Sequence 2410 Sequence 2411 Sequence 2412 Sequence 2413 Sequence 2414 Sequence 2415 Sequence 2416 Sequence 2417 Sequence 2418 Sequence 2418 Sequence 2419 Sequence 2419 Sequence 2419 Sequence 2419 Sequence 2420 Sequence 2420 Sequence 2421 Sequence 2420 Sequence 2421 Sequence 2421 Sequence 2421 Sequence 2422 Sequence 2423 Sequence 2424 Sequence 2424 Sequence 2424 Sequence 2425 Sequence 2426 Sequence 2427 Sequence 2427 Sequence 2428 Sequence 2429 Sequence 2430 Sequence 2430 Sequence 2430 Sequence 2430 Sequence 2430 Sequence 2430 Se	Sequence 2392	Al375141	Sequence 2444	Al970919
Sequence 2395 A/180137 Sequence 2447 A/459455 Sequence 2396 A/180137 Sequence 2448 A/476360 Sequence 2397 A/101742 Sequence 2449 A/1606272 Sequence 2398 A/1028097 Sequence 2450 A/1025262 Sequence 2399 A/8665185 Sequence 2451 H03785 Sequence 2400 A/071199 Sequence 2451 H03785 Sequence 2401 A/479962 Sequence 2453 A/082342 Sequence 2401 A/479962 Sequence 2453 A/082342 Sequence 2402 A/133862 Sequence 2454 A/186755 Sequence 2403 A/916145 Sequence 2455 A/086128 Sequence 2404 A/167677 Sequence 2456 A/086128 Sequence 2405 A/904265 Sequence 2457 A/086128 Sequence 2406 A/040284 Sequence 2459 A/086128 Sequence 2407 A/243143 Sequence 2459 A/086428 Sequence 2408 A/093458 Sequence 2459 A/086428 Sequence 2409 A/045881 Sequence 2460 A/045881 Sequence 2410 A/736613 Sequence 2462 A/287720 Sequence 2411 A/748064 Sequence 2463 A/086438 Sequence 2412 A/05253 Sequence 2465 A/058673 Sequence 2413 A/05404 Sequence 2463 A/086438 Sequence 2414 A/748064 Sequence 2465 A/058674 Sequence 2416 A/0437 Sequence 2466 A/036673 Sequence 2417 A/05663 Sequence 2466 A/036873 Sequence 2418 A/0437 Sequence 2468 A/038712 Sequence 2419 A/048064 Sequence 2468 A/038712 Sequence 2410 A/0470 Sequence 2468 A/058712 Sequence 2411 A/048064 Sequence 2468 A/058861 Sequence 2412 A/048064 Sequence 2466 A/038712 Sequence 2413 A/048064 Sequence 2466 A/038712 Sequence 2414 A/048064 Sequence 2468 A/038712 Sequence 2416 A/0487 Sequence 2476 A/0489129 Sequence 2417 A/048065 Sequence 2477 A/0489129 Sequence 2418 A/04037 Sequence 2478 A/0489129 Sequence 2419 A/04037 Sequence 2479 A/0489129 Sequence 2421 A/04037 Sequence 2479 A/0489129 Sequence 2422 A/048076 Sequence 2477 A/0489129 Sequence 2423 A/048676 Sequence 2477 A/0489129 Sequence 2424 A/0437 Sequence 2479 A/0499129 Sequence 2425 A/04966 Sequence 2479 A/0499129 Sequence 2426 A/04966 Sequence 2479 A/0499129 Sequence 2427 A/058676 Sequence 2489 A/0499129 Sequence 2428 A/04969 Sequence 2489 A/0499129 Sequence 2429 A/04999 Sequence 2489 A/049993 Sequence 2489 A/049993 Sequence 2489 A/049993 Sequence 2489 A/049993 Sequenc	Sequence 2393	N63784		
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Page 24 of 74

Table 4-1

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Page 25 of 74

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Table 4-1

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Sequence 2753	AA493551	Sequence 2805	AA120782
Sequence 2754	Al267652	Sequence 2806	AA129551
Sequenc∈ 2755	A1085554	Sequence 2807	AA215777

Page 27 of 74

Sequence 2808	AA150459	Sequence 2860	AA809784
Sequence 2809	AA436546	Sequence 2861	AA502798
Sequence 2810	A1609659	Sequence 2862	AA568207
Sequence 2811	AA031528	Sequence 2863	Al524414
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Sequence 2815	AA040694	Sequence 2867	A1762634
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Sequence 2818	AA513052	Sequence 2870	Al267162
Sequence 2819	AA427459	Sequence 2871	AA463533
Sequence 2820	AA044586	Sequence 2872	N89861
Sequence 2821	AA379449	Sequence 2873	AA206675
Sequence 2822	A1625404	Sequence 2874	AA418477
Sequence 2823	AA333300	Sequence 2875	AA639972
Sequence 2824	A1253300	Sequence 2876	Al620284
Sequence 2825	AI253424	Sequence 2877	AA515908
Sequence 2826	Al374628	Sequence 2878	AA133742
Sequence 2827	AA069845	Sequence 2879	H49828
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Sequence 2829	AA469199	Sequence 2881	AA505816
Sequence 2830	Al267416	Sequence 2882	AA121836
Sequence 2831	AA234135	Sequence 2883	R52729
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Sequence 2834	H78888	Sequence 2886	A1588087
Sequence 2835	T81369	Sequence 2887	AW152265
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Sequence 2837	AA812581	Sequence 2889	N24284
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Sequence 2857	A1469767	Sequence 2909	AW173278
Sequence 2858	R18975	Sequence 2910	AA296298
Sequence 2859	AA196978	Sequence 2911	Al312130
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Page 28 of 74

Sequence 2912	AA369786	Sequence 2964	AA425407
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Sequence 2914	AA206004	Sequence 2966	A1688573
Sequence 2915	AI627750	Sequenœ 2967	AA227579
Sequence 2916	AI340246	Sequence 2968	AA131834
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Sequence 2918	A1267492	Sequence 2970	AA575898
Sequence 2919	AI207515	Sequence 2971	AA195259
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Sequence 2922	AA151796	Sequence 2974	AA452200
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Sequence 2936	AA454921	Sequence 2988	Al478365
Sequence 2937	AA164983	Sequence 2989	AA890010
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Sequence 2939	AA321703	Sequence 2991	AA429336
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Sequence 2961	AA972598	Sequence 3013	AA304947
Sequence 2962	N28331	Sequence 3014	AA449544
Sequence 2963	R46295	Sequence 3015	AA583461
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Page 29 of 74

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Sequence 3022	AW167665		AA092316
Sequence 3023	AA573557	Sequence 3075	AA577598
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Sequence 3026	AA367948	Sequence 3078	AA442145
Sequence 3027	AA399265	Sequence 3079	AA680052
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Sequence 3067		Sequence 3119	AA826649
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Page 30 of 74

Sequence 3120	AA299140	Sequence 3172	J02642
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Sequence 3122	N36967	Sequence 3174	AB007972
Sequence 3123	AA400418	Sequence 3175	AF151837
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	M17323	Sequence 3219	AF109219
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Sequence 3170	D10495	Sequence 3221	L09235
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Page 31 of 74

Sequence 3224	L05425	<b>334</b>	AF070553
Sequence 3225	X68560	- · ·	AF032885
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Sequence 3227	D63486		875755
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Sequence 3251	U12596	Sequence 3303	S78203
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Sequence 3275	м 10905	Sequence 3327	AB032948
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Page 32 of 74

Table 4-1

Sequence 3328	L15702	Sequence 3380	E03569
Sequence 3329	AF017748	Sequence 3381	D28476
Sequence 3330	U14528	Sequence 3382	AF091263
Sequence 3331	AF098533	Sequence 3383	AB032981
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Sequence 3341	AF153608	Sequence 3393	J03537
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Sequence 3344	AF198358	Sequence 3396	AL049969
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Sequence 3357	AB033078	Sequence 3409	D50927
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Sequence 3360	AF070561	Sequence 3412	AF191018
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Sequence 3362	M69066	Sequence 3414	AB021288
Sequence 3363	AF057299	Sequence 3415	M97347
Sequence 3364	D38555	Sequence 3416	J05211
Sequence 3365	AL050198	Sequence 3417	AF131844
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Sequence 3378	J02853	Sequence 3430	S77601
Sequence 3379	AF061258	Sequence 3431	AL122076

Page 33 of 74

Sequence 3432	D50926	Sequence 3484	U53468
Sequence 3433	AF151836	Sequence 3485	M28372
Sequence 3434	U51990	Sequence 3486	AF058718
Sequence 3435	M17325	Sequence 3487	J03007
Sequence 3436	D50420	Sequence 3488	AL117430
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Sequence 3440	AL050186	Sequence 3492	AF035319
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Page 34 of 74

Sequence 353		Sequence 3588	AF100755
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Sequence 355	59 D31886	Sequence 3611	D14705
Sequence 356	60 AF084457	Sequence 3612	AF047442
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Sequence 356	33 X75535	Sequence 3615	D14043
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Sequence 358		Sequence 3635	AF107405
Sequence 358		Sequence 3636	D80001
Sequence 358	35 AL117392	Sequence 3637	M31627
Sequence 358		Sequence 3638	AF000652
Sequence 35	87 AB023173	Sequence 3639	M69238

Page 35 of 74

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Sequence 3640	AF070649	Sequence 3692	X41346
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Sequence 3643	M64572	Sequence 3695	V90283
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Sequence 3655	D87444	Sequence 3707	V62428
Sequence 3656	AB028948	Sequence 3708	V99733
Sequence 3657	AF022212	Sequence 3709	T72167
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Sequence 3663	AF038404	Sequence 3715	X26850
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Sequence 3667	X64838	Sequence 3719	Z33428
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Sequence 3669	J04615	Sequence 3721	Z15132
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Sequence 3676	AB018284	Sequence 3728	AC36117
Sequence 3677	D14812	Sequence 3729	AC36739
Sequence 3678	M34181	Sequence 3730	AC36708
Sequence 3679	Z13009	Sequence 3731	AC36079
Sequence 3680	L13773	Sequence 3732	AC36078
Sequence 3681	AF027824	Sequence 3733	AC36740
Sequence 3682	X82321	Sequence 3734	A21185
Sequence 3683	AB020694	Sequence 3735	AB000220
Sequence 3684	S69272	Sequence 3736	AB002303
Sequence 3685	AF070597	Sequence 3737	AB002369
Sequence 3686	AF091090	Sequence 3738	AB002387
Sequence 3687	U90548	Sequence 3739	AB007510
Sequence 3688	V62427	Sequence 3740	AB007884
Sequence 3689	V62430	Sequence 3741	AB007956
Sequence 3690	V35457	Sequence 3742	AB011004
Sequence 3691	V72211	Sequence 3743	AB014531

Page 36 of 74

Sequence 3744	AB014597	Sequence 3796	AF151103
Sequence 3745	AB016068	Sequence 3797	AF152306
Sequence 3746	AB018080	Sequence 3798	AF156965
Sequence 3747	AB018284	Sequence 3799	AF159056
Sequence 3748	AB023230	Sequence 3800	AF170583
Sequence 3749	AB032957	Sequence 3801	AF178574
Sequence 3750	AB032983	Sequence 3802	AF188745
Sequence 3751	AB033007	Sequence 3803	AJ001443
Sequence 3752	AB033078	Sequence 3804	AJ002744
Sequence 3753	AB033079	Sequence 3805	AJ004913
Sequence 3754	AB033091	Sequence 3806	AJ010069
Sequence 3755	AF000231	Sequence 3807	AJ223812
Sequence 3756	AF001212	Sequence 3808	AJ224875
Sequence 3757	AF007144	Sequence 3809	AL035304
Sequence 3758.	AF007216	Sequence 3810	AL049229
Sequence 3759	AF007217	Sequence 3811	AL049274
Sequence 3760	AF010233	Sequence 3812	AL049932
Sequence 3761	AF010312	Sequence 3813	AL049941
Sequence 3762	AF012126	Sequence 3814	AL049969
Sequence 3763	AF012549	Sequence 3815	AL050073
Sequence 3764	AF014955	Sequence 3816	AL080084
Sequence 3765	AF020202	Sequence 3817	AL110126
Sequence 3766	AF020736	Sequence 3818	AL117392
Sequence 3767	AF020761	Sequence 3819	AL117423
Sequence 3768	AF026166	Sequence 3820	AL117576
Sequence 3769	AF028832	Sequence 3821	AL117644
Sequence 3770	AF030403	Sequence 3822	AL117665
Sequence 3771	AF038404	Sequence 3823	D00039
Sequence 3772	AF039918	Sequence 3824	D13641
Sequence 3773	AF044956	Sequence 3825	D13866
Sequence 3774	AF047020	Sequence 3826	D14043
Sequence 3775	AF054838	Sequence 3827	D16562
Sequence 3776	AF060219	Sequence 3828	D29954
Sequence 3777	AF061258	Sequence 3829	D38305
Sequence 3778	AF065388	Sequence 3830	D80005
Sequence 3779	AF070553	Sequence 3831	D87742
Sequence 3780 Sequence 3781	AF071202 AF077037	Sequence 3832	D88208
Sequence 3782	AF077037	Sequence 3833	D90226
Sequence 3783	AF0778849	Sequence 3834	E01956
Sequence 3784	AF081281	Sequence 3835	E02628 E03569
Sequence 3785	AF084555	Sequence 3836 Sequence 3837	E03509 E07798
Sequence 3786	AF085845	Sequence 3838	
Sequence 3787	AF086477	Sequence 3839	J02923
Sequence 3788	AF092133	Sequence 3840	J04205 J04794
Sequence 3789	AF092565	Sequence 3841	
Sequence 3790	AF103907	Sequence 3842	J05192
Sequence 3791	AF132937	Sequence 3843	K01911
Sequence 3792	AF134159	Sequence 3844	L07033
Sequence 3793	AF144566	Sequence 3845	L38961 M10036
Sequence 3794	AF146277	•	M10036
Sequence 3795	AF147339	Sequence 3846	M12938
-04a0.00 0130	/11 1-71 005	Sequence 3847	M17885

Page 37 of 74

Sequence 3848	M20260	Sequence 3900	AA111855
Sequence 3849	M21895	Sequence 3901	AA113811
Sequence 3850	M24194	Sequence 3902	AA126472
Sequence 3851	M24902	Sequence 3903	AA135001
Sequence 3852	M26663	Sequence 3904	AA136221
Sequence 3853	M29366	Sequence 3905	AA136404
Sequence 3854	M30938	Sequence 3906	AA148087
Sequence 3855	M34840	Sequence 3907	AA148831
Sequence 3856	M36634	Sequence 3908	AA149579
Sequence 3857	M97934	Sequence 3909	AA149979
Sequence 3858	S82081	Sequence 3910	AA160149
Sequence 3859	U01923	Sequence 3911	AA167696
Sequence 3860	U02680	Sequence 3912	AA167829
Sequence 3861	U17104	Sequence 3913	AA171605
Sequence 3862	U33821	Sequence 3914	AA180845
Sequence 3863	U34252	Sequence 3915	AA187604
Sequence 3864	U36336	Sequence 3916	AA191536
Sequence 3865	· U50203	Sequence 3917	AA193086
Sequence 3866	U51990	Sequence 3918	AA195010
Sequence 3867	X04408	Sequence 3919	AA196978
Sequence 3868	X05332	Sequence 3920	AA203537
Sequence 3869	X15729	Sequence 3921	AA215793
Sequence 3870	X55654	Sequence 3922	AA216690
Sequence 3871	X67698	Sequence 3923	AA225294
Sequence 3872	X74331	Sequence 3924	AA226171
Sequence 3873	X75535	Sequence 3925	AA227000
Sequence 3874	X75861	Sequence 3926	AA228273
Sequence 3875	X80910	Sequence 3927	AA228940
Sequence 3876	X87241	Sequence 3928	AA229495
Sequence 3877	X94323	Sequence 3929	AA243809
Sequence 3878	X94910	Sequence 3930	AA250947
Sequence 3879	X98296	Sequence 3931	AA256117
Sequence 3880	Y00345	Sequence 3932	AA257955
Sequence 3881	Z11531	Sequence 3933	AA261819
Sequence 3882	Z26317	Sequence 3934	AA280647
Sequence 3883	Z46606	Sequence 3935	AA282286
Sequence 3884	Z48605	Sequence 3936	AA285341
Sequence 3885	AA001734	Sequence 3937	AA297647
Sequence 3886	AA002140	Sequence 3938	AA306876
Sequence 3887	AA026 <b>797</b>	Sequence 3939	AA307103
Sequence 3888	AA041467	Sequence 3940	AA308578
Sequence 3889	AA043109	Sequence 3941	AA310637
Sequence 3890	AA045803	Sequence 3942	AA311819
Sequence 3891	AA047482	Sequence 3943	AA312495
Sequence 3892	AA056437	Sequence 3944	AA312968
Sequence 3893	AA058899	Sequence 3945	AA314907
Sequence 3894	AA081003	Sequence 3946	AA315555
Sequence 3895	AA083026	Sequence 3947	AA316159
Sequence 3896	AA085835	Sequence 3948	AA326359
Sequence 3897	AA091277	Sequence 3949	AA326833
Sequence 3898	AA094000	Sequence 3950	AA334710
Sequence 3899	AA095049	Sequence 3951	AA336412

Table 4-1

Sequence 3952	AA345904	Sequence 4004	AA639786
Sequence 3953	AA346844	Sequence 4005	AA640474
Sequence 3954	AA355553	Sequence 4006	AA652478
Sequence 3955	AA359832	Sequence 4007	AA662733
Sequence 3956	AA368792	Sequence 4008	AA680052
Sequence 3957	AA375531	Sequence 4009	AA701667
Sequence 3958	AA399218	Sequence 4010	AA706224
Sequence 3959	AA399606	Sequence 4011	AA748031
Sequence 3960	AA418628	Sequence 4012	AA770006
Sequence 3961	AA418770	Sequence 4013	AA779670
Sequence 3962	AA420680	Sequence 4014	AA810905
Sequence 3963	AA424456	Sequence 4015	AA813941
Sequence 3964	AA426351	Sequence 4016	AA860839
Sequence 3965	AA430304	Sequence 4017	AA890010
Sequence 3966	AA436125	Sequence 4018	AA890326
Sequence 3967	AA437099	Sequence 4019	AA902625
Sequence 3968	AA437224	Sequence 4020	AA971532
Sequence 3969	AA442479	Sequence 4021	AA975275
Sequence 3970	AA447322	Sequence 4022	AA984384
Sequence 3971	AA455917	Sequence 4023	AA988520
Sequence 3972	AA460587	Sequence 4024	AA992583
Sequence 3973	AA460649	Sequence 4025	AF114027
Sequence 3974	AA461015	Sequence 4026	AF150208
Sequence 3975	AA484133	Sequence 4027	AI041116
Sequence 3976	AA469238	Sequence 4028	A1051210
Sequence 3977	AA471025	Sequence 4029	AI064967
Sequence 3978	AA480124	Sequence 4030	AI075338
Sequence 3979	AA488732	Sequence 4031	A1076042
Sequence 3980	AA489173	Sequence 4032	AI080267
Sequence 3981	AA492192	Sequence 4033	Al129991
Sequence 3982	AA497050	Sequence 4034	Al133562
Sequence 3983	AA501968	Sequence 4035	Al147472
Sequence 3984	AA502799	Sequence 4036	Al149363
Sequence 3985	AA502979	Sequence 4037	AI188850
Sequence 3986	AA516531	Sequence 4038	Al198311
Sequence 3987	AA523498	Sequence 4039	A1198937
Sequence 3988	AA523902	Sequence 4040	Al199681
Sequence 3989	AA524651	Sequence 4041	A1201248
Sequence 3990	AA528104	Sequence 4042	Al221390
Sequence 3991	AA531249	Sequence 4043	AI263876
Sequence 3992	AA533553	Sequence 4044	Al267162
Sequence 3993	AA541703	Sequence 4045	Al267185
Sequence 3994	AA543102	Sequence 4048	Al267216
Sequence 3995	AA548472	Sequence 4047	Al267254
Sequence 3996	AA548722	Sequence 4048	Al267282
Sequence 3997	AA569813	Sequence 4049	Al278643
Sequence 3998	AA570171	Sequence 4050	Al300562
Sequence 3999	AA576425	Sequence 4051	Al336326
Sequence 4000	AA578773	Sequence 4052	Al361256
Sequence 4001	AA580771	Sequence 4053	Al367759
Sequence 4002	AA608967	Sequence 4054	Al431801
Sequence 4003	AA635590	Sequence 4055	AI472067
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Page 39 of 74

Table 4-1

Sequence 4056	AI547327	Sequence 4108	H54240
Sequence 4057	AI557182	Sequence 4109	H78517
Sequence 4058	AI557225	Sequence 4110	H78888
Sequence 4059	Al55 <b>7235</b>	Sequence 4111	H79639
Sequence 4060	A1557277	Sequence 4112	N23545
Sequence 4061	AI564487	Sequence 4113	N27218
Sequence 4062	A1620427	Sequence 4114	N58818
Sequence 4063	A1627356	Sequence 4115	N62448
Sequence 4064	A1630306	Sequence 4116	R35529
Sequence 4065	Al630357	Sequence 4117	R55535
Sequence 4066	A1630362	Sequence 4118	R72738
Sequence 4067	A1633646	Sequence 4119	T08676
Sequence 4068	Al656122	Sequence 4120	T79397
Sequence 4069	AI658850	Sequence 4121	T79514
Sequence 4070	AI682937	Sequence 4122	W15195
Sequence 4071	AI686139	Sequence 4123	W19757
Sequence 4072	A16947 <b>6</b> 7	Sequence 4124	W25081
Sequence 4073	Al741213	Sequence 4125	W56069
Sequence 4074	Al743852	Sequence 4126	W67735
Sequence 4075	A1755085	Sequence 4127	W79779
Sequence 4076	A1766083	Sequence 4128	W88856
Sequence 4077	AI767692 I	Sequence 4129	T99948
Sequence 4078	A1799891	Sequence 4130	V54589
Sequence 4079	A1825408	Sequence 4131	V58628
Sequence 4080	AI826504	Sequence 4132	V63195
Sequence 4081	AI912057	Sequence 4133	X03841
Sequence 4082	Al914133	Sequence 4134	X04318
Sequence 4083	A1925735	Sequence 4135	X26850
Sequence 4084	Al949242	Sequence 4136	X27262
Sequence 4085	AI953152	Sequence 4137	X27278
Sequence 4086	Al968401	Sequence 4138	X39870
Sequence 4087	A1970381	Sequence 4139	X97676
Sequence 4088	AL036483	Sequence 4140	Z15277
Sequence 4089	AL042404	Sequence 4141	<b>Z27248</b>
Sequence 4090	AL043046	Sequence 4142	Z33526
Sequence 4091	AL047415	Sequence 4143	Z41241
Sequence 4092	AL047881	Sequence 4144	AC34465
Sequence 4093	AW022756	Sequence 4145	AC36078
Sequence 4094	AW044142	Sequence 4146	AC36079
Sequence 4095	AW068579	Sequence 4147	AC36099
Sequence 4096	AW149817	Sequence 4148	AC36175
Sequence 4097	AW160731	Sequence 4149	AC36180
Sequence 4098	AW163485	Sequence 4150	AC36201
Sequence 4099	AW176022	Sequence 4151	AC36457
Sequence 4100	AW179322	Sequence 4152	AC36740
Sequence 4101	C06419	Sequence 4153	AB020866
Sequence 4102	H00112	Sequence 4154	AC002040
Sequence 4103	H03785	Sequence 4155	AC002064
Sequence 4104	H06944	Sequence 4156	AC003991
Sequence 4105	H13891	Sequence 4157	AC004656
Sequence 4106	H15654	Sequence 4158	AC005154
Sequence 4107	H17741	Sequence 4159	AC005538

Page 40 of 74

Sequence 4160	AC005831	Sequence 4212	AA057490
Sequence 4161	AC007052	Sequence 4213	AA057861
Sequence 4162	AC007686	Sequence 4214	AA062731
Sequence 4163	AF061258	Sequence 4215	AA062859
Sequence 4164	AL034399	Sequence 4216	AA070349
Sequence 4165	AL035666	Sequence 4217	AA074142
Sequence 4166	AL049821	Sequence 4218	AA074729
Sequence 4167	AP000527	Sequence 4219	AA074869
Sequence 4168	AP000528	Sequence 4220	AA075133
Sequence 4169	AP000529	Sequence 4221	AA075459
Sequence 4170	G16768	Sequence 4222	AA081007
Sequence 4171	S65371	Sequence 4223	AA081355
Sequence 4172	U66083	Sequence 4224	AA082685
Sequence 4173	X70394	Sequence 4225	AA083479
Sequence 4174	Z94388	Sequence 4226	AA084987
Sequence 4175	AW235635	Sequence 4227	AA085483
Sequence 4176	AI383637	Sequence 4228	AA088372
Sequence 4177	AC005480	Sequence 4229	AA088770
Sequence 4178	AL031003	Sequence 4230	AA090106
Sequence 4179	AB020866	Sequence 4231	AA093276
Sequence 4180	S65371	Sequence 4232	AA094656
Sequence 4181	AA001673	Sequence 4233	AA096121
Sequence 4182	AA002131	Sequence 4234	AA099976
Sequence 4183	AA007370	Sequence 4235	AA100852
Sequence 4184	AA009611	Sequence 4236	AA100876
Sequence 4185	AA010689	Sequence 4237	AA112310
Sequence 4186	AA010981	Sequence 4238	AA114062
Sequence 4187	AA021608	Sequence 4239	AA114854
Sequence 4188	AA024581	Sequence 4240	AA114967
Sequence 4189	AA026399	Sequence 4241	AA115021
Sequence 4190	AA029143	Sequence 4242	AA115492
Sequence 4191	AA029567	Sequence 4243	AA120816
Sequence 4192	AA029889	Sequence 4244	AA121127
Sequence 4193	AA031360	Sequence 4245	AA121419
Sequence 4194	AA032221	Sequence 4246	AA122237
Sequence 4195	AA035001	Sequence 4247	AA122245
Sequence 4196	AA035773	Sequence 4248	AA125817
Sequence 4197	AA037425	Sequence 4249	AA126121
Sequence 4198	AA037620	Sequence 4250	AA126729
Sequence 4199	AA039807	Sequence 4251	AA1268 <b>0</b> 9
Sequence 4200	AA039998	Sequence 4252	AA126921
Sequence 4201	AA040180	Sequence 4253	AA127029
Sequence 4202	AA040710	Sequence 4254	AA127714
Sequence 4203	AA044118	Sequence 4255	AA128387
Sequence 4204	AA044586	Sequence 4256	AA128442
Sequence 4205	AA046083	Sequence 4257	AA129615
Sequence 4206	AA047432	Sequence 4258	AA129652
Sequence 4207	AA053265	Sequence 4259	AA129781
Sequence 4208	AA053932	Sequence 4260	AA130162
Sequence 4209	AA056345	Sequence 4261	AA130515
Sequence 4210	AA057063	Sequence 4262	AA130529
Sequence 4211	AA057270	Sequence 4263	AA130928

Page 41 of 74

Sequence 4264	AA131313	Sequence 4316	AA191134
Sequence 4265	AA131812	Sequence 4317	AA191552
Sequence 4266	AA131926	Sequence 4318	AA193346
Sequence 4267	AA132689	Sequence 4319	AA194517
Sequence 4268	AA132844	Sequence 4320	AA194860
Sequence 4269	AA133030	Sequence 4321	AA195010
Sequence 4270	AA133361	Sequence 4322	AA195044
Sequence 4271	AA133652	Sequence 4323	AA195063
Sequence 4272	AA134022	Sequence 4324	AA195145
Sequence 4273	AA134203 -	Sequence 4325	AA196558
Sequence 4274	AA135870	Sequence 4326	AA199802
Sequence 4275	AA136215	Sequence 4327	AA203592
Sequence 4276	AA137165	Sequence 4328	AA204701
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Sequence 4279	AA146582	Sequence 4331	AA205870
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Sequence 4285	AA149979	Sequence 4337	AA223306
Sequence 4286	AA150891	Sequence 4338	AA224124
Sequence 4287	AA151018	Sequence 4339	AA225865
Sequence 4288	AA151506	Sequence 4340	AA226137
Sequence 4289	AA156092	Sequence 4341	AA226171
Sequence 4290	AA156879	Sequence 4342	AA227506
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Sequence 4296	AA167150	Sequence 4348	AA229993
Sequence 4297	AA167674	Sequence 4349	AA232691
Sequence 4298	AA169374	Sequence 4350	AA233689
Sequence 4299	AA169794	Sequence 4351	AA233868
Sequence 4300	AA171510	Sequence 4352	AA234130
Sequence 4301	AA171872	Sequence 4353	AA236057
Sequence 4302	AA171883	Sequence 4354	AA236323
Sequence 4303	AA173363	Sequence 4355	AA243361
Sequence 4304	AA173997	Sequence 4356	AA247816
Sequence 4305	AA176331	Sequence 4357	AA251303
Sequence 4306	AA177127	Sequence 4358	AA251644
Sequence 4307	AA180137	Sequence 4359	AA252355
Sequence 4308	AA186346	Sequence 4360	AA255695
Sequence 4309	AA186399	Sequence 4361	AA255717
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Sequence 4312	AA188045	Sequence 4364	AA262296
Sequence 4313	AA188052	Sequence 4365	AA279347
Sequence 4314	AA188169	Sequence 4366	AA279949
Sequence 4315	AA190561	Sequence 4367	AA280694
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Page 42 of 74

Table 4-1

Sequence 43	368	AA280735	Sequence 4420	AA344743
Sequence 4	369	AA283704	Sequence 4421	AA345624
Sequence 4	370	AA284161	Sequence 4422	AA349118
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Sequence 4	372	AA284555	Sequence 4424	AA353766
Sequence 4	373 .	AA287751	Sequence 4425	AA356682
Sequence 4	374	AA287961	Sequence 4426	AA357811
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Sequence 4	376	AA291756	Sequence 4428	AA361403
Sequence 4	377 .	AA292692	Sequence 4429	AA366865
Sequence 4	378	AA292993	Sequence 4430	AA367471
Sequence 4	379	AA296846	Sequence 4431	AA368520
Sequence 4	380	AA297434	Sequence 4432	AA370337
Sequence 4	381	AA297883	Sequence 4433	AA370352
Sequence 4	382	AA298489	Sequence 4434	AA371017
Sequence 4	383	AA298786	Sequence 4435	AA371265
Sequence 4		AA298807	Sequence 4436	AA371626
Sequence 4	385	AA299830	Sequence 4437	AA375853
Sequence 4		AA300065	Sequence 4438	AA375988
Sequence 4	387	AA302852	Sequence 4439	AA376870
Sequence 4		AA302919	Sequence 4440	AA393480
Sequence 4	389	AA306206	Sequence 4441	AA394298
Sequence 4		AA306888	Sequence 4442	AA398686
Sequence 4		AA307209	Sequence 4443	AA399133
Sequence 4		AA307247	Sequence 4444	AA401339
Sequence 4		AA307789	Sequence 4445	AA401480
Sequence 4		AA309445	Sequence 4446	AA402095
Sequence 4		AA310663	Sequence 4447	AA403051
Sequence 4		AA310739	Sequence 4448	AA405663
Sequence 4		AA312756	Sequence 4449	AA406475
Sequence 4		AA313223	Sequence 4450	AA410580
Sequence 4		AA314022	Sequence 4451	AA418796
Sequence 4		AA314127	Sequence 4452	AA420845
Sequence 4		AA314188	Sequence 4453	AA421021
Sequence 4		AA314673	Sequence 4454	AA421562
Sequence 4		AA314872 AA315095	Sequence 4455	AA425394
Sequence 4			Sequence 4456	AA426351
Sequence 4		AA315259	Sequence 4457	AA427477
Sequence 4 Sequence 4		AA315805 AA315838	Sequence 4458	AA429912
Sequence 4		AA316627	Sequence 4459	AA431677
Sequence 4		AA318745	Sequence 4460	AA436377
Sequence 4	– –	AA320004	Sequence 4461	AA437224
Sequence 4		AA322115	Sequence 4462	AA441885
Sequence 4		AA325145	Sequence 4463 Sequence 4464	AA441918
Sequence 4		AA325826	•	AA441959
Sequence 4		AA330090	Sequence 4465	AA442287
Sequence 4		AA333526	Sequence 4466	AA442372
Sequence 4		AA334457	Sequence 4467	AA442829
Sequence 4		AA334754	Sequence 4468	AA443750
Sequence 4		AA335699	Sequence 4469	AA443790
Sequence 4		AA336586	Sequence 4470	AA443946
- 24401106 4	7713	/V 1000000	Sequence 4471	AA444074

Page 43 of 74

Sequence 4472	AA445955	Sequence 4524	AA508588
Sequence 4473	AA446116	Sequence 4525	AA508889
Sequence 4474	AA446149	Sequence 4528	AA508902
Sequence 4475	AA447322	Sequence 4527	AA512974
Sequence 4476	AA447948	Sequence 4528	AA513177
Sequence 4477	AA448036	Sequence 4529	AA513688
Sequence 4478	AA448531	Sequence 4530	AA515857
Sequence 4479	AA449604	Sequence 4531	AA516531
Sequence 4480	AA451633	Sequence 4532	AA522481
Sequence 4481	AA453612	Sequence 4533	AA522856
Sequence 4482	AA454208	Sequence 4534	AA522867
Sequence 4483	AA458514	Sequence 4535	AA523252
Sequence 4484	AA459617	Sequence 4536	AA523464
Sequence 4485	AA460120	Sequence 4537	AA523484
Sequence 4486	AA461082	Sequence 4538	AA523938
Sequence 4487	AA461548	Sequence 4539	AA524812
Sequence 4488	AA463198	Sequence 4540	AA526886
Sequence 4489	AA464133	Sequence 4541	AA527730
Sequence 4490	AA465193	Sequence 4542	AA527805
Sequence 4491	AA465258	Sequence 4543	AA528104
Sequence 4492	AA465293	Sequence 4544	AA531563
Sequence 4493	AA465387	Sequence 4545	AA533553
Sequence 4494	AA468839	Sequence 4546	AA533772
Sequence 4495	AA469151	Sequence 4547	AA534171
Sequence 4496	AA478514	Sequence 4548	AA53441B
Sequence 4497	AA479327	Sequence 4549	AA534540
Sequence 4498	AA481001	Sequence 4550	AA534933
Sequence 4499	AA481303	Sequence 4551	AA535837
Sequence 4500	AA482654	Sequence 4552	AA541677
Sequence 4501	AA485861	Sequence 4553	AA541678
Sequence 4502	AA485906	Sequence 4554	AA541808
Sequence 4503	AA485996	Sequence 4555	AA542906
Sequence 4504	AA486628	Sequence 4556	AA548238
Sequence 4505	AA487261	Sequence 4557	AA551146
Sequence 4506	AA488227	Sequence 4558	AA565420
Sequence 4507	AA488744	Sequence 4559	AA565882
Sequence 4508	AA488809	Sequence 4560	AA569130
Sequence 4509	AA489020	Sequence 4561	AA569809
Sequence 4510	AA489433	Sequence 4562	AA573824
Sequence 4511	AA492042	Sequence 4563	AA574288
Sequence 4512	AA492280	Sequence 4564	AA574397
Sequence 4513	AA493512	Sequence 4565	AA576425
Sequence 4514	AA495759	Sequence 4566	AA578773
Sequence 4515	AA495938	Sequence 4567	AA580900
Sequence 4516	AA502979	Sequence 4568	AA581117
Sequence 4517 Sequence 4518	AA503182	Sequence 4569	AA582252
	AA503215	Sequence 4570	AA582596
Sequence 4519	AA503265	Sequence 4571	AA588474
Sequence 4520	AA503731	Sequence 4572	AA600998
Sequence 4521 Sequence 4522	AA504238	Sequence 4573	AA602430
Sequence 4523	AA504340	Sequence 4574	AA602794
Oequence 4023	AA505198	Sequence 4575	AA603334

Page 44 of 74

Table 4-1

Sequence 4576	AA604474	Sequence 4628	AA773641
Sequence 4577	AA612666	Sequence 4629	AA773787
Sequence 4578	AA612829	Sequence 4630	AA776585
Sequence 4579	AA612874	Sequence 4631	AA779176
Sequence 4580	AA618277	Sequence 4632	AA780591
Sequence 4581	AA620750	Sequence 4633	AA782138
Sequence 4582	AA625951	Sequence 4634	AA782644
Sequence 4583	AA626509	Sequence 4635	AA805865
Sequence 4584	AA628417	Sequence 4636	AA806808
Sequence 4585	AA629027	. Sequence 4637	AA811365
Sequence 4586	AA631868	Sequence 4638	AA813108
Sequence 4587	AA639059	Sequence 4639	AA825612
Sequence 4588	AA639174	Sequence 4640	AA828708
Sequence 4589	AA639791	Sequence 4641	AA829473
Sequence 4590	AA640687	Sequence 4642	AA833543
Sequence 4591	AA641636	Sequence 4643	AA836233
Sequence 4592	AA641769	Sequence 4644	AA843242
Sequence 4593	AA641810	Sequence 4645	AA843414
Sequence 4594	AA642383	Sequence 4646	AA843663
Sequence 4595	AA643506	Sequence 4647	AA845340
Sequence 4596	AA644237	Sequence 4648	AA847380
Sequence 4597	AA650230	Sequence 4649	AA854293
Sequence 4598	AA652206	Sequence 4650	AA857509
Sequence 4599	AA652478	Sequence 4651	AA860541
Sequence 4600	AA662733	Sequence 4652	AA860986
Sequence 4601	AA664567	Sequence 4653	AA865873
Sequence 4602	AA665507	Sequence 4654	AA868720
Sequence 4603	AA679314	Sequence 4655	AA873182
Sequence 4604	AA679547	Sequence 4656	AA873291
Sequence 4605	AA687216	Sequence 4657	AA873828
Sequence 4606	AA702230	Sequence 4658	AA876668
Sequence 4607	AA706388	Sequence 4659	AA885770
Sequence 4608	AA707607	Sequence 4660	AA888014
Sequence 4609	AA708016	Sequence 4661	AA890327
Sequence 4610	AA708690	Sequence 4662	AA902127
Sequence 4611	AA714481	Sequence 4663	AA902721
Sequence 4612	AA722204	Sequence 4664	AA902964
Sequence 4613	AA724553	Sequence 4665	AA905139
Sequence 4614	AA737949	Sequence 4666	AA911245
Sequence 4615	AA743680	Sequence 4667	AA916753
Sequence 4616	AA744367	Sequence 4668	AA931228
Sequence 4617	AA744771	Sequence 4669	AA931237
Sequence 4618	AA747515	Sequence 4670	AA931583
Sequence 4619	AA747772	Sequence 4671	AA932114
Sequence 4620	AA748847	Sequence 4672	AA935526
Sequence 4621	AA761640	Sequence 4673	AA935560
Sequence 4622	AA764991	Sequence 4674	AA936262
Sequence 4623	AA765884	Sequence 4675	AA937551
Sequence 4624	AA765974	Sequence 4676	AA938717
Sequence 4625	AA766644	Sequence 4677	AA946873
Sequence 4626	AA769055	Sequence 4678	AA953892
Sequence 4627	AA773072	Sequence 4679	AA954118

Page 45 of 74

Table 4-1

Sequence 4680	AA968756	Sequence 4732	Al142257
Sequence 4681	AA974540	Sequence 4733	Al144488
Sequence 4682	AA975275	Sequence 4734	Al148552
Sequence 4683	AA977172	Sequence 4735	Al148676
Sequence 4684	AA985327	Sequence 4736	Al148933
Sequence 4685	AA987997	Sequence 4737	Al159768
Sequence 4686	AA988125	Sequence 4738	A1174675
Sequence 4687	AA988520	Sequence 4739	Al198311
Sequence 4688	AA995981	Sequence 4740	Al198577
Sequence 4689	AI004252	Sequence 4741	Al198986
Sequence 4690	AI015053	Sequence 4742	Al199094
Sequence 4691	Al017442	Sequence 4743	Al199681
Sequence 4692	AI017996	Sequence 4744	Al201483
Sequence 4693	AI023770	Sequence 4745	Al203141
Sequence 4694	A1024638	Sequence 4746	Al203395
Sequence 4695	A1024799	Sequence 4747	A1207650
Sequence 4696	AI026122	Sequence 4748	Al214272
Sequence 4697	Al027643	Sequence 4749	Al214784
Sequence 4698	AI032759	Sequence 4750	Al219278
Sequence 4699	AI033912	Sequence 4751	Al220149
Sequence 4700	AI034131	Sequence 4752	Al221713
Sequence 4701	AI038167	Sequence 4753	Al245327
Sequence 4702	AI038415	Sequence 4754	AJ247078
Sequence 4703	Al040598	Sequence 4755	Al248736
Sequence 4704	AI041616	Sequence 4758	Al250092
Sequence 4705	AI042153	Sequence 4757	Al253335
Sequence 4706	AI051210	Sequence 4758	Al253348
Sequence 4707	AI051376	Sequence 4759	AI253379
Sequence 4708	AI052453	Sequence 4760	Al267185
Sequence 4709	Al061420	Sequence 4761	AI267282
Sequence 4710	AI064763	Sequence 4762	Al267454
Sequence 4711	AI066518	Sequence 4763	Al267502
Sequence 4712	A1074969	Sequence 4764	Al267574
Sequence 4713	A1075935	Sequence 4765	A1268724
Sequence 4714	A1077439	Sequence 4766	A1274786
Sequence 4715	AI078772	Sequence 4767	Al278659
Sequence 4716	A1080439	Sequence 4768	AI283489
Sequence 4717	A1080476	Sequence 4769	Al288142
Sequence 4718	AI084101	Sequence 4770	A1289059
Sequence 4719	AI085977	Sequence 4771	Al299945
Sequence 4720	A1086377	Sequence 4772	AI300489
Sequence 4721	A1087987	Sequence 4773	Al301532
Sequence 4722	AI088201	Sequence 4774	AI308800
Sequence 4723	A1091190	Sequence 4775	Al310465
Sequence 4724	A1093553	Sequence 4776	Al312542
Sequence 4725	Al125642	Sequence 4777	Al313387
Sequence 4726	Al125688	Sequence 4778	Al332588
Sequence 4727	Al126233	Sequence 4779	Al333116
Sequence 4728	Al126700	Sequence 4780	AI338338
Sequence 4729	AI127167	Sequence 4781	Al339458
Sequence 4730	Al129321	Sequence 4782	Al343692
Sequence 4731	Al141040	Sequence 4783	Al346021
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Page 46 of 74

Sequence 4784	Al346657	Sequence 4836	AI682088
Sequence 4785	A1357472	Sequence 4837	A1682937
Sequence 4786	A1358723	Sequence 4838	Al686779
Sequence 4787	Al366381	Sequence 4839	A1693302
Sequence 4788	Al366549	Sequence 4840	Al694199
Sequence 4789	A13 <b>73289</b>	Sequence 4841	Al694596
Sequence 4790	Al394646	Sequence 4842	A1696992
Sequence 4791	Al400410	Sequence 4843	A1697849
Sequence 4792	Al420172	Sequence 4844	A1734266
Sequence 4793	Al421741	Sequence 4845	Al740796
Sequence 4794	Al423541	Sequence 4846	AI743852
Sequence 4795	A1432462	Sequence 4847	A1749547
Sequence 4796	Al434150	Sequence 4848	Al753784
Sequence 4797	Al452485	Sequence 4849	Al754013
Sequence 4798	Al457964	Sequence 4850	AI755085
Sequence 4799	Al458673	Sequence 4851	AI762085
Sequence 4800	A1460230	Sequence 4852	AI769392
Sequence 4801	Al467858	Sequence 4853	AI769478
Sequence 4802	A1475476	Sequence 4854	AI791370
Sequence 4803	A1478365	Sequence 4855	AI791906
Sequence 4804	AI493851	Sequence 4856	Al791998
Sequence 4805	AI521984	Sequence 4857	AI801810
Sequence 4806	AI524826	Sequence 4858	AI805379
Sequence 4807	AI525102	Sequence 4859	A1806044
Sequence 4808	AI546840	Sequence 4860	AI813772
Sequence 4809	AI557059	Sequence 4861	AI816102
Sequence 4810	AI557112	Sequence 4862	AI828602
Sequence 4811	AI557118	Sequence 4863	AI829784
Sequence 4812	A1557225	Sequence 4864	AI861997
Sequence 4813	AI557243	Sequence 4865	A1864927
Sequence 4814	AI557363	Sequence 4866	AI871027
Sequence 4815	A1557495	Sequence 4867	AI871745
Sequence 4816	A1559299	Sequence 4868	AI879995
Sequence 4817	A1559903	Sequence 4869	AIB84963
Sequence 4818	AI560870	Sequence 4870	Al915264
Sequence 4819	A1564487	Sequence 4871	AI923011
Sequence 4820	A1589533	Sequence 4872	AI935704
Sequence 4821	A1608684	Sequence 4873	AI935710
Sequence 4822	A1608968	Sequence 4874	A1936472
Sequence 4823	Al610412	Sequence 4875	AI948822
Sequence 4824	Al619498	Sequence 4876	AI954585
Sequence 4825	Al620796	Sequence 4877	AI973022
Sequence 4826	A1627553	Sequence 4878	Al973109
Sequence 4827	A1635310	Sequence 4879	A1985122
Sequence 4828	AI635658	Sequence 4880	AL036594
Sequence 4829	Al635667	Sequence 4881	AL036801
Sequence 4830	AI659898	Sequence 4882	AL037456
Sequence 4831	Al660356	Sequence 4883	AL037895
Sequence 4832	A1660919	Sequence 4884	AL038821
Sequence 4833	Al671394	Sequence 4885	AL039175
Sequence 4834	AI679619	Sequence 4886	AL039235
Sequence 4835	AI680874	Sequence 4887	AL039293

Page 47 of 74

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Sequence 4888	AL039349	Sequence 4940	H57382
Sequence 4889	AL040204	Sequence 4941	H65423
Sequence 4890	AL043702	Sequence 4942	H69529
Sequence 4891	AL044985	Sequence 4943	H83215
Sequence 4892	AL119715	Sequence 4944	H87419
Sequence 4893	AL120972	Sequence 4945	H95535
Sequence 4894	AW019951	Sequence 4946	H96053
Sequence 4895	AW020930	Sequence 4947	H96527
Sequence 4896	AW021628	Sequence 4948	N34611
Sequence 4897	AW022112	Sequence 4949	N35555
Sequence 4898	AW028521	Sequence 4950	N42986
Sequence 4899	AW052140	Sequence 4951	N47044
Sequence 4900	AW055209	Sequence 4952	N49471
Sequence 4901	AW058600	Sequence 4953	N49534
Sequence 4902	AW080631	Sequence 4954	N52554
Sequence 4903	AW080845	Sequence 4955	N76215
Sequence 4904	AW088727	Sequence 4956	N92498
Sequence 4905	AW131770	Sequence 4957	N94829
Sequence 4906	AW134672	Sequence 4958	N98569
Sequence 4907	AW135187	Sequence 4959	N98672
Sequence 4908	AW135430	Sequence 4960	R14008
Sequence 4909	AW135666	Sequence 4961	R17476
Sequence 4910	AW136951	Sequence 4962	R19498
Sequence 4911	AW138413	Sequence 4963	R19761
Sequence 4912	AW140049	Sequence 4964	R24400
Sequence 4913	AW148750	Sequence 4965	R25577
Sequence 4914	AW150770	Sequence 4966	R33762
Sequence 4915	AW160363	Sequence 4967	R34284
Sequence 4916	AW178825	Sequence 4968	R35797
Sequence 4917	C05816	Sequence 4969	R49734
Sequence 4918	C06196	Sequence 4970	R52387
Sequence 4919	C06419	Sequence 4971	R59846
Sequence 4920	D58694	Sequence 4972	R61601
Sequence 4921	F07635	Sequence 4973	R63823
Sequence 4922	F12493	Sequence 4974	R70570
Sequence 4923	F13272	Sequence 4975	R75754
Sequence 4924	F22763	Sequence 4976	T29959
Sequence 4925	F33993	Sequence 4977	T34760
Sequence 4926	H00742	Sequence 4978	T48624
Sequence 4927	H02725	Sequence 4979	T66793
Sequence 4928	H03785	Sequence 4980	T70123
Sequence 4929	H08511	Sequence 4981	T72034
Sequence 4930	H09882	Sequence 4982	T83709
Sequence 4931	H12923	Sequence 4983	T86219
Sequence 4932	H13008	Sequence 4984	T89530
Sequence 4933	H13406	Sequence 4985	T99386
Sequence 4934	H13891	Sequence 4986	W03386
Sequence 4935	H20744	Sequence 4987	W04191
Sequence 4936	H25211	Sequence 4988	W05240
Sequence 4937	H49331	Sequence 4989	W05402
Sequence 4938	H49828	Sequence 4990	W06897
Sequence 4939	H54240	Sequence 4991	W19970
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Page 48 of 74

Table 4-1

Sequence 4992	W21894	Sequence 5044	AB018340
Sequence 4993	W22330	Sequence 5045	AB018346
Sequence 4994	W24228	Sequence 5046	AB019568
Sequence 4995	W26413	Sequence 5047	AB020631
Sequence 4996	W30686	Sequence 5048	AB020643
Sequence 4997	W31819	Sequence 5049	AB020657
Sequence 4998	W37614	Sequence 5050	AB020694
Sequence 4999	W44623	Sequence 5051	AB020724
Sequence 5000	W44911	Sequence 5052	AB021663
Sequence 5001	W44954	Sequence 5053	AB021981
Sequence 5002	W56395	Sequence 5054	AB023157
Sequence 5003	W60565	Sequence 5055	AB023173
Sequence 5004	W61042	Sequence 5056	AB023177
Sequence 5005	W72973	Sequence 5057	AB023227
Sequence 5006	W74046	Sequence 5058	AB023230
Sequence 5007	W80673	Sequence 5059	AB024334
Sequence 5008	W96214	Sequence 5060	AB024435
Sequence 5009	Z43304	Sequence 5061	AB028257
Sequence 5010	A06805	Sequence 5062	AB028624
Sequence 5011	A06846	Sequence 5063	AB028973
Sequence 5012	A06919	Sequence 5064	AB028995
Sequence 5013	A16794	Sequence 5065	AB029036
Sequence 5014	AB000888	Sequence 5066	AB032948
Sequence 5015	AB001106	Sequence 5067	AB032983
Sequence 5016	AB002303	Sequence 5068	AB033042
Sequence 5017	AB002330	Sequence 5069	AB033071
Sequence 5018	AB002347	Sequence 5070	AB033075
Sequence 5019	AB002365	Sequence 5071	AB033076
Sequence 5020	AB002369	Sequence 5072	AB033078
Sequence 5021	AB002370	Sequence 5073	AB033079
Sequence 5022	AB002382	Sequence 5074	AB033080
Sequence 5023	AB004047	Sequence 5075	AB033092
Sequence 5024	AB004857	Sequence 5076	AB033899
Sequence 5025	AB007510	Sequence 5077	AD001528
Sequence 5026	AB007896	Sequence 5078	AF000652
Sequence 5027	AB007899	Sequence 5079	AF000982
Sequence 5028	AB007916	Sequence 5080	AF000983
Sequence 5029	AB007931	Sequence 5081	AF001902
Sequence 5030	AB007934	Sequence 5082	AF002668
Sequence 5031	AB007956	Sequence 5083	AF002697
Sequence 5032	AB007969	Sequence 5084	AF006088
Sequence 5033	AB008226	Sequence 5085	AF007216
Sequence 5034	AB009285	Sequence 5086	AF007791
Sequence 5035	AB011004	Sequence 5087	AF008442
Sequence 5036	AB011079	Sequence 5088	AF010233
Sequence 5037	AB011148	Sequence 5089	AF013759
Sequence 5038	AB011159	Sequence 5090	AF015553
Sequence 5039	AB014560	Sequence 5091	AF016371
Sequence 5040	AB015317	Sequence 5092	AF019214
Sequence 5041	AB016068	Sequence 5093	AF019226
Sequence 5042	AB018266	Sequence 5094	AF020736
Sequence 5043	AB018337	Sequence 5095	AF020797

Page 49 of 74

Sequence 5096	AF021351	Sequence 5148	AF073299
Sequence 5097	AF026166	Sequence 5149	AF074606
Sequence 5098	AF026246	Sequence 5150	AF075061
Sequence 5099	AF027824	Sequence 5151	AF077030
Sequence 5100	AF028593	Sequence 5152	AF077202
Sequence 5101	AF028832	Sequence 5153	AF078845
Sequence 5102	AF029669	Sequence 5154	AF078855
Sequence 5103	AF030162	Sequence 5155	AF078860
Sequence 5104	AF033095	Sequence 5156	AF081282
Sequence 5105	AF035285	Sequence 5157	AF083106
Sequence 5106	AF035293	Sequence 5158	AF085224
Sequence 5107	AF035316	Sequence 5159	AF085845
Sequence 5108	AF037447	Sequence 5160	AF085871
Sequence 5109	AF038187	Sequence 5181	AF085896
Sequence 5110	AF038451	Sequence 5162	AF086172
Sequence 5111	AF038452	Sequence 5163	AF086245
Sequence 5112	AF038955	Sequence 5164	AF086336
Sequence 5113	AF039023	Sequence 5165	AF087980
Sequence 5114	AF039918	Sequence 5166	AF091084
Sequence 5115	AF042181	Sequence 5167	AF092092
Sequence 5116	AF044953	Sequence 5168	AF092138
Sequence 5117	AF044956	Sequence 5169	AF097514
Sequence 5118	AF047020	Sequence 5170	AF097725
Sequence 5119	AF047470	Sequence 5171	AF099137
Sequence 5120	AF047472	Sequence 5172	AF100756
Sequence 5121	AF048731	Sequence 5173	AF101074
Sequence 5122	AF051334	Sequence 5174	AF102846
Sequence 5123	AF052107	Sequence 5175	AF103907
Sequence 5124	AF052154	Sequence 5176	AF106681
Sequence 5125	AF053535	Sequence 5177	AF106862
Sequence 5126	AF053641	Sequence 5178	AF109219
Sequence 5127	AF054181	Sequence 5179	AF110377
Sequence 5128	AF054186	Sequence 5180	AF111106
Sequence 5129	AF054187	Sequence 5181	AF118838
Sequence 5130	AF054284	Sequence 5182	AF124141
Sequence 5131	AF054838	Sequence 5183	AF124438
Sequence 5132	AF054990	Sequence 5184	AF124598
Sequence 5133	AF057160	Sequence 5185	AF126245
Sequence 5134	AF061258	Sequence 5186	AF131743
Sequence 5135	AF061739	Sequence 5187	AF131775
Sequence 5136	AF062323	Sequence 5188	AF131831
Sequence 5137	AF064257	Sequence 5189	AF131856
Sequence 5138	AF064768	Sequence 5190	AF131857
Sequence 5139	AF068235	Sequence 5191	AF132000
Sequence 5140	AF070523	Sequence 5192	AF132940
Sequence 5141	AF070556	Sequence 5193	AF132965
Sequence 5142	AF070561	Sequence 5194	AF133123
Sequence 5143	AF070621	Sequence 5195	AF135421
Sequence 5144	AF070646	Sequence 5196	AF144103
Sequence 5145	AF070659	Sequence 5197	AF144566
Sequence 5146	AF070661	Sequence 5198	AF146277
Sequence 5147	AF071202	Sequence 5199	AF147331

Page 50 of 74

Table 4-1

Sequence 5200	AF151103	Sequence 5252	AL079298
Sequence 5201	AF151109	Sequence 5253	AL080059
Sequence 5202	AF151840	Sequence 5254	AL080084
Sequence 5203	AF151851	Sequence 5255	AL080118
Sequence 5204	AF151871	Sequence 5256	AL080192
Sequence 5205	AF151B75	Sequence 5257	AL080209
Sequence 5206	AF151897	Sequence 5258	AL080212
Sequence 5207	AF152097	Sequence 5259	AL110126
Sequence 5208	AF153608	Sequence 5260	AL110177
Sequence 5209	AF155099	Sequence 5261	AL110183
Sequence 5210	AF156098	Sequence 5262	AL110207
Sequence 5211	AF159056	Sequence 5263	AL110222
Sequence 5212	AF159615	Sequence 5264	AL110233
Sequence 5213	AF168956	Sequence 5265	AL110241
Sequence 5214	AF171877	Sequence 5266	AL117392
Sequence 5215	AF174605	Sequence 5267	AL117616
Sequence 5216	AF176555	Sequence 5268	AL122121
Sequence 5217	AF176574	Sequence 5269	AL133108
Sequence 5218	AF176700	Sequence 5270	AL133117
Sequence 5219	AF188746	Sequence 5271	D10040
Sequence 5220	AF188747	Sequence 5272	D13286
Sequence 5221	AF201077	Sequence 5273	D13315
Sequence 5222	AJ001381	Sequence 5274	D13388
Sequence 5223	AJ001443	Sequence 5275	D13627
Sequence 5224	AJ002744	Sequence 5276	D13630
Sequence 5225	AJ002955	Sequence 5277	D13641
Sequence 5226	AJ004913	Sequence 5278	D13866
Sequence 5227	AJ005821	Sequence 5279	D13900
Sequence 5228	AJ007398	Sequence 5280	D14446
Sequence 5229	AJ011007 ·	Sequence 5281	D14530
Sequence 5230	AJ012499	Sequence 5282	D14657
Sequence 5231	AJ223352	Sequence 5283	D14658
Sequence 5232	AJ223353	Sequence 5284	D14696
Sequence 5233	AJ224326	Sequence 5285	D14710
Sequence 5234	AJ249248	Sequence 5286	D14812
Sequence 5235	AL049266	Sequence 5287	D14878
Sequence 5236	AL049447	Sequence 5288	D16307
Sequence 5237	AL049471	Sequence 5289	D16960
Sequence 5238	AL049705	Sequence 5290	D17039
Sequence 5239	AL049932	Sequence 5291	D17130
Sequence 5240	AL049969	Sequence 5292	D17400
Sequence 5241	AL049996	Sequence 5293	D21163
Sequence 5242	AL050013	Sequence 5294	D21260
Sequence 5243	AL050018	Sequence 5295	D25283
Sequence 5244	AL050035	Sequence 5296	D25542
Sequence 5245	AL050064	Sequence 5297	D26129
Sequence 5246	AL050071	Sequence 5298	D26600
Sequence 5247	AL050161	Sequence 5299	D29640
Sequence 5248	AL050186	Sequence 5300	D29954
Sequence 5249	AL050204	Sequence 5301	D31885
Sequence 5250	AL050268	Sequence 5302	D37991
Sequence 5251	AL079288	Sequence 5303	D38305

Page 51 of 74

## Table 4-1

Sequence 5304	D38551	Sequence 5356	K00558
Sequence 5305	D38552	Sequence 5357	K01911
Sequence 5306	D42047	Sequence 5358	L01100
Sequence 5307	D42138	Sequence 5359	L02426
Sequence 5308	D45887	Sequence 5360	L05425
Sequence 5309	D49396	Sequence 5361	L06132
Sequence 5310	D50487	Sequence 5362	L06133
Sequence 5311	D50525	Sequence 5363	L06850
Sequence 5312	D50926	Sequence 5364	L07633
Sequence 5313	D63480	Sequence 5365	L07758
Sequence 5314	D64015	Sequence 5366	L08441
Sequence 5315	D78333	Sequence 5367	L10284
Sequence 5316	D79996	Sequence 5368	L10678
Sequence 5317	D80005	Sequence 5369	L11566
Sequence 5318	D80009	Seguence 5370	L11667
Sequence 5319	D82345	Sequence 5371	L12686
Sequence 5320	D85777	Sequence 5372	L13977
Sequence 5321	D86322	Sequence 5373	L15203
Sequence 5322	D86971	Sequence 5374	L17128
Sequence 5323	D87443	Sequence 5375	L29158
Sequence 5324	D87444	Sequence 5376	L34839
Sequence 5325	D87453	Sequence 5377	L34840
Sequence 5326	D87742	Sequence 5378	L35249
Sequence 5327	D87845	Sequence 5379	L37080
Sequence 5328	D88674	Sequence 5380	L38608
Sequence 5329	D89053	Sequence 5381	L38951
Sequence 5330	D89667	Sequence 5382	L38961
Sequence 5331	D89675	Sequence 5383	L47647
Sequence 5332	D90373	Sequence 5384	M10036
Sequence 5333	E01574	Sequence 5385	M10119
Sequence 5334	E01650	Sequence 5386	M11058
Sequence 5335	E02628	Sequence 5387	M11353
Sequence 5336	E03414	Sequence 5388	M11887
Sequence 5337	E03569	Sequence 5389	M11949
Sequence 5338	E05957	Sequence 5390	M15885
Sequence 5339	E06721	Sequence 5391	M15887
Sequence 5340	J02642	Sequence 5392	M15990
Sequence 5341	J02888	Sequence 5393	M16247
Sequence 5342	J02923	Sequence 5394	M16768
Sequence 5343	J02966	Sequence 5395	M16804
Sequence 5344	J03007	Sequence 5396	M16827
Sequence 5345	J03015	Sequence 5397	M17323
Sequence 5346	J03143	Sequence 5398	M19961
Sequence 5347	J03248	Sequence 5399	M20372
Sequence 5348	J03503	Sequence 5400	M20471
Sequence 5349	J03575	Sequence 5401	M21154
Sequence 5350	J03592	Sequence 5402	M21895
Sequence 5351	J03934	Sequence 5403	M22382
Sequence 5352	J04162	Sequence 5404	M22590
Sequence 5353	J04183	Sequence 5405	M23161
Sequence 5354	J05021	Sequence 5408	M23613
Sequence 5355	J05480	Sequence 5407	M24194

Page 52 of 74

Sequence 5408	M24902	Sequence 5460	U05877
Sequençe 5409	M26325	Sequence 5461	U07151
Sequence 5410	M26481	Sequence 5462	U07919
Sequence 5411	M26663	Sequence 5463	U10485
Sequence 5412	M27504	Sequence 5464	U12465
Sequence 5413	M27691	Sequence 5465	U12789
Sequence 5414	M28211	Sequence 5466	U13616
Sequence 5415	M28372	Sequence 5467	U14750
Sequence 5416	M28526	Sequence 5468	U14967
Sequence 5417	M29064	Sequence 5469	U14968
Sequence 5418	M29872	Sequence 5470	U14970
Sequence 5419	M31630	Sequence 5471	U15008
Sequence 5420	M33197	Sequence 5472	U17040
Sequence 5421	M34175	Sequence 5473	U18197
Sequence 5422	M34181	Sequence 5474	U18914
Sequence 5423	M34840	Sequence 5475	U19177
Sequence 5424	M35252	Sequence 5476	U25182
Sequence 5425	M36634	Sequence 5477	U25789
Sequence 5426	.M37400	Sequence 5478	U27143
Sequence 5427	M37583	Sequence 5479	U28249
Sequence 5428	M38690	Sequence 5480	U30826
Sequence 5429	M55409	Sequence 5481	U31383
Sequence 5430	M55536	Sequence 5482	U33760
Sequence 5431	M58581	Sequence 5483	U33821
Sequence 5432	M59465	Sequence 5484	U34252
Sequence 5433	M62401	Sequence 5485	U37230
Sequence 5434	M63573	Sequence 5486	U37518
Sequence 5435	M65131	Sequence 5487	U39400
Sequence 5436	M69226	Sequence 5488	U40272
Sequence 5437	M69238	Sequence 5489	U42404
Sequence 5438	M74091	Sequence 5490	U43899
Sequence 5439	M74509	Sequence 5491	U47414
Sequence 5440	M74777	Sequence 5492	U54558
Sequence 5441	M81757	Sequence 5493	U57052
Sequence 5442	M82919	Sequence 5494	U57057
Sequence 5443	M84443	Sequence 5495	U58855
Sequence 5444	M86917	Sequence 5496	U59919
Sequence 5445	M90054	Sequence 5497	U60205
Sequence 5446	M90309	Sequence 5498	U61232
Sequence 5447	M90516	Sequence 5499	U62740
Sequence 5448	M94856	Sequence 5500	U63542
Sequence 5449	M95724	Sequence 5501	U68140
Sequence 5450	M96803	Sequence 5502	U68758
Sequence 5451	S50223	Sequence 5503	U70063
Sequence 5452	S57235	Sequence 5504	U70322
Sequence 5453	S59049	Sequence 5505	U70451
Sequence 5454	S64650	Sequence 5506	U72761
Sequence 5455	S68015	Sequence 5507	U75272
Sequence 5456	S69272	Sequence 5508	U75679
Sequence 5457	S73498	Sequence 5509	U785 <b>7</b> 5
Sequence 5458	U02680	Sequence 5510	U80034
Sequence 5459	U05291	Sequence 5511	U83115

Page 53 of 74

### Table 4-1

Sequence 5512	U856 <b>58</b>	Sequence 5564	X97065
Sequence 5513	U87166	Sequence 5565	X97675
Sequence 5514	U90915	Sequence 5566	X98263
Sequence 5515	U96759	Sequence 5567	Y00282
Sequence 5516	V00478	Sequence 5568	Y00345
Sequence 5517	X02530	Sequence 5569	Y08991
Sequence 5518	X04098	Sequence 5570	Y09615
Sequence 5519	X05332	Sequence 6571	Y09836
Sequence 5520	X05345	Sequence 5572	Y12065
Sequence 5521	X06272	Sequence 5573	Y12860
Sequence 5522	X06617	Sequence 5574	Y13286
Sequence 5523	X12433	Sequence 5575	Y15286
Sequence 5524	X12791	Sequence 5576	Y16132
Sequence 5525	X13425	Sequence 5577	Y16241
Sequence 5526	X13482	Sequence 5578	Y16414
Sequence 5527	X13585	Sequence 5579	Y18314
Sequence 5528	X13839	Sequence 5580	Z21507
Sequence 5529	X15187	Sequence 5581	Z22658
Sequence 5530	X15729	Sequence 5582	Z36531
Sequence 5531	X52882	Sequence 5583	Z37986
Sequence 5532	X53605	Sequence 5584	Z47087
Sequence 5533	X53793	Sequence 5585	Z47553
Sequence 5534	X54199	Sequence 5586	N91825
Sequence 5535	X54473	Sequence 5587	Q92813
Sequence 5536	X56932	Sequence 5588	T47520
Sequence 5537	X61970	Sequence 5589	T59274
Sequence 5538	X62534	Sequence 5590	T84934
Sequence 5539	X63753	. Sequence 5591	V04715
Sequence 5540	X64330	Sequence 5592	V07572
Sequence 5541	X64838	Sequence 5593	V44867
Sequence 5542	X6 <b>6276</b>	Sequence 5594	V58694
Sequence 5543	X66785	Sequence 5595	V62427
Sequence 5544	X67098	Sequence 5596	V63172
Sequence 5545	X73114	Sequence 5597	V70354
Sequence 5546	X73608	Sequence 5598	X03841
Sequence 5547	X76770	Sequence 5599	X24060
Sequence 5548	X76771	Sequence 5600	X26850
Sequence 5549	X77196	Sequence 5601	X27262
Sequence 5550	X7862 <b>7</b>	Sequence 5602	X29296
Sequence 5551	X79201	Sequence 5603	X37405
Sequence 5552	X81198	Sequence 5604	X37486
Sequence 5553	X81695	Sequence 5605	X40518
Sequence 5554	X82157	Sequence 5606	X41115
Sequence 5555	X84908	Sequence 5607	X52268
Sequence 5556	X85337	Sequence 5608	X87388
Sequence 5557	X85372	Sequence 5609	X89852
Sequence 5558	X85373	Sequence 5610	Z00428
Sequence 5559	X86098	Sequence 5611	Z13254
Sequence 5560	X87176	Sequence 5612	Z14392
Sequence 5561	X87949	Sequence 5613	Z15132
Sequence 5562	· X94323	Sequence 5614	AC29430
Sequence 5563	X95384	Sequence 5615	A06919

Page 54 of 74

Table 4-1

Sequence 5616	A06925	Sequence 5668	AJ132637
Sequence 5617	AB0 <b>02370</b>	Sequence 5669	AJ250042
Sequence 5618	AB002375	Sequence 5670	AL049932
Sequence 5619	AB0 <b>06534</b>	Sequence 5671	AL117499
Sequence 5620	AB007510	Sequence 5672	D30655
Sequence 5621	AB007899	Sequence 5673	D38551
Sequence 5622	AB011079	Sequence 5674	D42044
Sequence 5623	AB014458	Sequence 5675	D49396
Sequence 5624	AB015639	Sequence 5676	D50371
Sequence 5625	AB016068	Sequence 5677	D50525
Sequence 5626	AB018276	Sequence 5678	E02628
Sequence 5627	AB018313	Sequence 5679	J02959
Sequence 5628	AB020646	Sequence 5680	J03007
Sequence 5629	AB0 <b>20693</b>	Sequence 5681	K01144
Sequence 5630	AB020694	Sequence 5682	K01911
Sequence 5631	AB023173	Sequence 5683	L09235
Sequence 5632	AB023182	Sequence 5684	L10678
Sequence 5633	AB023227	Sequence 5685	L12136
Sequence 5634	AB033114	Sequence 5686	L15203
Sequence 5635	AF007791	Sequence 5687	L34839
Sequence 5636	AF015040	Sequence 5688	M10119
Sequence 5637	AF016371	Sequence 5689	M13231
Sequence 5638	AF019632	Sequence 5690	M15885
Sequence 5639	AF020797	Sequence 5691	M16553
Sequence 5640	AF044195	Sequence 5692	M16768
Sequence 5641	AF047020	Sequence 5693	M17885
Sequence 5642	AF052642	Sequence 5694	M24902
Sequence 5643	AF061258	Sequence 5695	M26663
Sequence 5644	AF067008	Sequence 5696	M74777
Sequence 5645	AF070525	Sequence 5697	M97935
Sequence 5646	AF070561	Sequence 5698	S69272
Sequence 5647	AF071202	Sequence 5699	S69738
Sequence 5648	AF077202	Sequence 5700	S75755
Sequence 5649	AF078854	Sequence 5701	U07919
Sequence 5650	AF085845	Sequence 5702	U21936
Sequence 5651	AF093097	Sequence 5703	U40272
Sequence 5652	AF100756	Sequence 5704	U42404
Sequence 5653	AF100759	Sequence 5705	U79751
Sequence 5654	AF103907	Sequence 5706	U96759
Sequence 5655	AF104914	Sequence 5707	X05606
Sequence 5656	AF126008 AF126736	Sequence 5708	X06617
Sequence 5657		Sequence 5709	X12791
Sequence 5658 Sequence 5659	AF131856 AF132000	Sequence 5710	X15187
Sequence 5660	AF146277	Sequence 5711	X52174
Sequence 5661		Sequence 5712	X78627
Sequence 5662	AF151103 AF151893	Sequence 5713	X85372
Sequence 5663	AF151693 AF153612	Sequence 5714	X92845
Sequence 5664	AF159056	Sequence 5715	X94323
Sequence 5665	AF188746	Sequence 5716	Y00345
Sequence 5666	AF191298	Sequence 5717	Z11793
Sequence 5667	AF 19 1298 AJ002744	Sequence 5718	AA026715
Sequence 5007	AUUU2144	Sequence 5719	AA033901

Page 55 of 74

Sequence 5720	AA035189	Sequence 5772	AA485217
Sequence 5721	AA043802	Sequence 5773	AA485935
Sequence 5722	AA081708	Sequence 5774	AA489433
Sequence 5723	AA083482	Sequence 5775	AA491041
Sequence 5724	AA092533	Sequence 5776	AA491249
Sequence 5725	AA094656	Sequence 5777	AA505076
Sequence 5726	AA100486	Sequence 5778	AA506953
Sequence 5727	AA128739	Sequence 5779	AA507804
Sequence 5728	AA133307	Sequence 5780	AA515143
Sequence 5729	AA135001	Sequence 5781	AA533057
Sequence 5730	AA148729	Sequence 5782	AA541587
Sequence 5731	AA156369	Sequence 5783	AA573762
Sequence 5732	AA156971	Sequence 5784	AA578209
Sequence 5733	AA171510	Sequence 5785	AA578773
Sequence 5734	AA172266	Sequence 5786	AA581222
Sequence 5735	AA179315	Sequence 5787	AA602522
Sequence 5736	AA181102	Sequence 5788	AA602957
Sequence 5737	AA182502	Sequence 5789	AA609384
Sequence 5738	AA189150	Sequence 5790	AA644495
Sequence 5739	AA203252	Sequence 5791	AA652478
Sequence 5740	AA205496	Sequence 5792	AA659693
Sequence 5741	AA224124	Sequence 5793	AA683246
Sequence 5742	AA225115	Sequence 5794	AA687216
Sequence 5743	AA236774	Sequence 5795	AA708003
Sequence 5744	AA249195	Sequence 5796	AA759202
Sequence 5745	AA258750	Sequence 5797	AA760894
Sequence 5746	AA262783	Sequence 5798	AA779728
Sequence 5747	AA278475	Sequence 5799	AA806808
Sequence 5748	AA291041	Sequence 5800	AA831278
Sequence 5749	AA292642	Sequence 5801	AA861930
Sequence 5750	AA298475	Sequence 5802	AA878195
Sequence 5751	AA299404	Sequence 5803	AA913138
Sequence 5752	AA305599	Sequence 5804	AA935560
Sequence 5753	AA306883	Sequence 5805	AA947278
Sequence 5754	AA321319	Sequence 5806	AA953892
Sequence 5755	AA333358	Sequence 5807	AA974999
Sequence 5756	AA339960	Sequence 5808	AA975275
Sequence 5757	AA370352	Sequence 5809	AA984134
Sequence 5758	AA370501	Sequence 5810	AI018034
Sequence 5759	AA371265	Sequence 5811	AI018625
Sequence 5760	AA393313	Sequence 5812	A1049995
Sequence 5761	AA400692	Sequence 5813	A1084651
Sequence 5762	AA422007	Sequence 5814	AI090226
Sequence 5763	AA433848	Sequence 5815	Al125642
Sequence 5764	AA441885	Sequence 5816	AI126233
Sequence 5765	AA447401	Sequence 5817	AI133406
Sequence 5766	AA454557	Sequence 5818	Al133727
Sequence 5767	AA455668	Sequence 5819	AI200578
Sequence 5768	AA468839	Sequence 5820	Al282318
Sequence 5769	AA478518	Sequence 5821	Al300489
Sequence 5770	AA479342	Sequence 5822	Al309401
Sequence 5771	AA479860	Sequence 5823	Al422907
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Page 56 of 74

Sequence 58	24 A	453358	Sequence 5	876	U <b>73642</b>
Sequence 58	25 AI	460230	Sequence 5	877	AC005880
Sequence 58	26 A	479021	Sequence 5	878	AF178030
Sequence 58	27 AI	480214	Sequence 5	879	AF072847
Sequence 58	28 AI	497759	Sequence 5	0888	AL121790
Sequence 58	29 A	557112	Sequence 5	5881	AC007279
Sequence 58	30 AI	581840	Sequence 5	882	AA001116
Sequence 58	131 AI	632108	Sequence 5	5883	AA001734
Sequence 58	132 AI	678832	Sequence 5	5884	AA001794
Sequence 58	133 AI	734266	Sequence 5	885	AA004802
Sequence 58	134 AI	755085	Sequence 5	5886	AA009818
Sequence 58	35 A	796571	Sequence 5	887	AA009878
Sequence 58	36 A	815454	Sequence 5	888	AA010174
Sequence 58	137 A	864040	Sequence 5	588 <del>9</del>	AA010682
Sequence 58	38 A	884963	Sequence 5	5890	AA010981
Sequence 58	39 AI	L037316	Sequence 5	891	AA011070
Sequence 58	140 AI	L037732	Sequence 5	5892	AA013090
Sequence 58	141 AI	L039293	Sequence 5	5893	AA017474
Sequence 58	42 A	L039599	Sequence 5	58 <b>9</b> 4	AA018591
Sequence 58	43 A	W014145	Sequence 5	5895	AA019781
Sequence 58	344 A	W020934	Sequence 5	5896	AA019874
Sequence 58	45 A	W085924	Sequence 5	5897	AA022636
Sequence 58	46 A	W135430	Sequence 5	5898	AA022937
Sequence 58	47 A	W173228	Sequence 5	5899	AA024796
Sequence 58	148 D	58694	Sequence 5	5900	AA025089
Sequence 58	49. H	51866	Sequence 5	5901	AA025968
Sequence 58	50 N	40017	Sequence 5	5902	AA026983
Sequence 58	851 R	22748	Sequence 5	5903	AA029039
Sequence 58	352 R	24400	Sequence 5	5904	AA029070
Sequence 58	153 R	49734	Sequence 5	5905	AA029143
Sequence 58		61436	Sequence !	5906	AA029276
Sequence 58		78584	Sequence 5	5907	AA029876
Sequence 58		81995	Sequence 5	5908	AA032220
Sequence 58		99386	Sequence !	5909	AA033651
Sequence 58		<i>[</i> 24708	Sequence (	5910	AA033907
Sequence 58		59274	Sequence 5	5911	AA034979
Sequence 58		06834	Sequence :	5912	AA035790
Sequence 58		24060	Sequence 5	5913	AA037210
Sequence 58		27262	Sequence (	5914	AA037483
Sequence 58		52268	Sequence !	5915	AA037886
Sequence 58		13254	Sequence 8	5916	AA039381
Sequence 58		C36079	Sequence !	5917	AA040810
Sequence 58		C003991	Sequence !	5918	AA043137
Sequence 58		C007275	Sequence (		AA043159
Sequence 58		P000502	Sequence !	5920	AA043415
Sequence 58		C007239	Sequence :	5921	AA044239
Sequence 58		C000065	Sequence (	5922	AA044941
Sequence 58		C005881	Sequence :	5923	AA045147
Sequence 58		C004982	Sequence :	5924	AA045410
Sequence 58		C003991	Sequence :		AA045803
Sequence 58		C004022	Sequence :	5926	AA046848
Sequence 58	3/5 A	C004706	Sequence !	5927	AA046973

Page 57 of 74

Table 4-1

Sequence 5928	AA046981	Sequence 5980	AA099110
Sequence 5929	AA053909	Sequence 5981	AA099976
Sequence 5930	AA054492	Sequence 5982	AA100852
Sequence 5931	AA056060	Sequence 5983	AA101777
Sequence 5932	AA056358	Sequence 5984	AA112626
Sequence 5933	AA056365	Sequence 5985	AA113403
Sequence 5934	AA056396	Sequence 5986	AA115021
Sequence 5935	AA056528	Sequence 5987	AA115297
Sequence 5936	AA057071	Sequence 5988	AA115838
Sequence 5937	AA057290	Sequence 5989	AA115933
Sequence 5938	AA058398	Sequence 5990	AA121419
Sequence 5939	AA058899	Sequence 5991	AA126029
Sequence 5940	AA058902	Sequence 5992	AA126742
Sequence 5941	AA062583	Sequence 5993	AA128114
Sequence 5942	AA062994	Sequence 5994	AA128647
Sequence 5943	AA063247	Sequence 5995	AA128668
Sequence 5944	AA069367	Sequence 5996	AA129772
Sequence 5945	AA069561	Sequence 5997	AA129964
Sequence 5946	AA071165	Sequence 5998	AA130207
Sequence 5947	AA074086	Sequence 5999	AA130349
Sequence 5948	AA074183	Sequence 6000	AA131227
Sequence 5949	AA074445	- Sequence 6001	AA131252
Sequence 5950	AA074509	Sequence 6002	AA131284
Sequence 5951	AA075460	Sequence 6003	AA131793
Sequence 5952	AA076295	Sequence 6004	AA132510
Sequence 5953	AA081007	Sequence 6005	AA132792
Sequence 5954	AA081355	Sequence 6006	AA133068
Sequence 5955	AA082209	Sequence 6007	AA133268
Sequence 5956	AA082572	Sequence 6008	AA133788
Sequence 5957	AA083410	Sequence 6009	AA134701
Sequence 5958	AA083411	Sequence 6010	AA135870
Sequence 5959	AA083482	Sequence 6011	AA137165
Sequence 5960	AA083510	Sequence 6012	AA143242
Sequence 5961	AA083772	Sequence 6013	AA143523
Sequence 5962	AA083836	Sequence 6014	AA143548
Sequence 5963	AA084221	Sequence 6015	AA146582
Sequence 5964	AA084698	Sequence 6016	AA146900
Sequence 5965	AA085480	Sequence 6017	AA147608
Sequence 5966	AA086007	Sequence 6018	AA147812
Sequence 5967	AA088679	Sequence 6019	AA148019
Sequence 5968	AA088758	Sequence 6020	AA148269
Sequence 5969	AA090106	Sequence 6021	AA148569
Sequence 5970	AA090203	Sequence 6022	AA149490
Sequence 5971	AA090548	Sequence 6023	AA149543
Sequence 5972	AA091275	Sequence 6024	AA149551
Sequence 5973	AA092060	Sequence 6025	AA149579
Sequence 5974	AA092343	Sequence 6026	AA149852
Sequence 5975	AA092532	Sequence 6027	AA150480
Sequence 5976	AA092533	Sequence 6028	AA150650
Sequence 5977	AA093935	Sequence 6029	AA150780
Sequence 5978	AA094331	Sequence 6030	AA150891
Sequence 5979	AA094656	Sequence 6031	AA151596
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Page 58 of 74

Table 4-1

Sequence 6032	AA152073	Sequence 6084	AA203206
Sequence 6033	AA152266	Sequence 6085	AA203285
Sequence 6034	AA152440	Sequence 6086	AA203637
Sequence 6035	AA155828	Sequence 6087	AA205033
Sequence 6036	AA155865	Sequence 6088	AA205647
Sequence 6037	AA156369	Sequence 6089	AA205857
Sequence 6038	AA157632	Sequence 6090	AA206769
Sequence 6039	AA159272	Sequence 6091	AA210760
Sequence 6040	AA159647	Sequence 6092	AA214668
Sequence 6041	AA159771	Sequence 6093	AA214688
Sequence 6042	AA160677	Sequence 6094	AA218788
Sequence 6043	AA165027	Sequence 6095	AA223249
Sequence 6044	AA165082	Sequence 6096	AA223433
Sequence 6045	AA165165	Sequence 6097	AA223956
Sequence 6046	AA165632	Sequence 6098	AA224124
Sequence 6047	AA166707	Sequence 6099	AA224994
Sequence 6048	AA167399	Sequence 6100	AA225115
Sequence 6049	AA169507	Sequence 6101	AA225294
Sequence 6050	AA171510	Sequence 6102	AA225488
Sequence 6051	AA173391	Sequence 6103	AA226171
Sequence 6052	AA173559	Sequence 6104	AA226234
Sequence 6053	AA177050	Sequence 6105	AA226321
Sequence 6054	AA179187	Sequence 6106	AA228273
Sequence 6055	AA179462	Sequence 6107	AA228431
Sequence 6056	AA179487	Sequence 6108	AA228822
Sequence 6057	AA180137	Sequence 6109	AA228940
Sequence 6058	AA180154	Sequence 6110	AA229495
Sequence 6059	AA181618	Sequence 6111	AA229611
Sequence 6060	AA182457	Sequence 6112	AA232691
Sequence 6061	AA186538	Sequence 6113	AA232700
Sequence 6062	AA187928	Sequence 6114	AA233073
Sequence 6063	AA187998	Sequence 6115	AA233088
Sequence 6064	AA188052	Sequence 6116	AA233131
Sequence 6065	AA188312	Sequence 6117	AA234073
Sequence 6066	AA188396	Sequence 6118	AA234092
Sequence 6067	AA188875	Sequence 6119	AA234990
Sequence 6068	AA189150	Sequence 6120	AA235914
Sequence 6069	AA190529	Sequence 6121	AA236320
Sequence 6070	AA190661	Sequence 6122	AA242891
Sequence 6071	AA190843	Sequence 6123	AA243143
Sequence 6072	AA190890	Sequence 6124	AA243696
Sequence 6073	AA191045	Sequence 6125	AA244003
Sequence 6074	AA191719	Sequence 6126	AA244116
Sequence 6075	AA191721	Sequence 6127	AA244207
Sequence 6076	AA192597	Sequence 6128	AA250748
Sequence 6077	AA194062	Sequence 6129	AA251178
Sequence 6078	AA194988	Sequence 6130	AA251578
Sequence 6079	AA196001	Sequence 6131	AA251644
Sequence 6080	AA196392	Sequence 6132	AA252849
Sequence 6081	AA196978	Sequence 6133	AA253453
Sequence 6082	AA199658	Sequence 6134	AA255761
Sequence 6083	AA203172	Sequence 6135	AA256330
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Page 59 of 74

Sequence 6136	AA257973	Sequence 6188	AA312059
Sequence 6137	AA258357	Sequence 6189	AA312591
Sequence 6138	AA258513	Sequence 6190	AA312756
Sequence 6139	AA261990	Sequence 6191	AA313057
Sequence 6140	AA262513	Sequence 6192	AA313244
Sequence 6141	AA263153	Sequence 6193	AA313653
Sequence 6142	AA278520	Sequence 6194	AA313979
Sequence 6143	AA279517	Sequence 6195	AA314659
Sequence 6144	AA280917	Sequence 6196	AA314810
Sequence 6145	AA284615	Sequence 6197	AA315016
Sequence 6146	AA286953	Sequence 6198	AA316411
Sequence 6147	AA287511	Sequence 6199	AA317068
Sequence 6148	AA291325	Sequence 6200	AA317496
Sequence 6149	AA291363	Sequence 6201	AA317976
Sequence 6150	AA291602	Sequence 6202	AA320004
Sequence 6151	AA291764	Sequence 6203	AA320723
Sequence 6152	AA292638	Sequence 6204	AA322540
Sequence 6153	AA293028	Sequence 6205	AA324478
Sequence 6154	AA293629	Sequence 6206	AA326306
Sequence 6155	AA295182	Sequence 6207	AA327105
Sequence 6156	AA295788	Seguence 6208	AA327469
Sequence 6157	AA296846	Sequence 6209	AA328770
Sequence 6158	AA297941	Sequence 6210	AA329242
Sequence 6159	AA298085	Sequence 6211	AA329770
Sequence 6160	AA298104	Sequence 6212	AA329937
Sequence 6161	AA298294	Sequence 6213	AA330123
Sequence 6162	AA298464	Sequence 6214	AA332083
Sequence 6163	AA298489	Sequence 6215	AA332593
Sequence 6164	AA300065	Sequence 6216	AA332716
Sequence 6165	AA302457	Sequence 6217	AA333090
Sequence 6166	AA305358	Sequence 6218	AA333526
Sequence 6167	AA305627	Sequence 6219	AA333765
Sequence 6168	AA305824	Sequence 6220	AA334344
Sequence 6169	AA305895	Sequence 6221	AA338215
Sequence 6170	AA305909	Sequence 6222	AA340635
Sequence 6171	AA306264	Sequence 6223	AA344625
Sequence 6172	AA306413	Sequence 6224	AA349232
Sequence 6173	AA306480	Sequence 6225	AA351297
Sequence 6174	AA306573	Sequence 6226	AA352938
Sequence 6175	AA306812	Sequence 6227	AA354872
Sequence 6176	AA306958	Sequence 6228	AA358790
Sequence 6177	AA306982	Sequence 6229	AA359838
Sequence 6178	AA306983	Sequence 6230	AA360190
Sequence 6179	AA307513	Sequence 6231	AA360310
Sequence 6180	AA307697	Sequence 6232	AA364171
Sequence 6181	AA308091	Sequence 6233	AA364844
Sequence 6182	AA309909	Sequence 6234	AA365620
Sequence 6183	AA310173	Sequence 6235	AA366995
Sequence 6184	AA310807	Sequence 6236	AA367471
Sequence 6185	AA311028	Sequence 6237	AA370110
Sequence 6186	AA311905	Sequence 6238	AA370222
Sequence 6187	AA311978	Sequence 6239	AA370352
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Page 60 of 74

Table 4-1

Sequence 6240	AA370507	Se	quence 6292	AA442976
Sequence 6241	AA370620	Se	equence 6293	AA443024
Sequence 6242	AA371017	_	quence 6294	AA443082
Sequence 6243	AA371157	Se	quence 6295	AA443242
Sequence 6244	AA371265	Se	quence 6296	AA444093
Sequence 6245	AA374020	Se	equence 6297	AA446298
Sequence 6246	AA374198	Se	quence 6298	AA446910
Sequence 6247	AA374814	Se	quence 6299	AA447608
Sequence 6248	AA375927	Se	quence 6300	AA447948
Sequence 6249	AA376034	Se	equence 6301	AA448795
Sequence 6250	AA378933	· Se	quence 6302	AA448882
Sequence 6251	AA379448	Se	quence 6303	AA449520
Sequence 6252	AA397832	Se	quence 6304	AA452012
Sequence 6253	AA397953	Se	quence 6305	AA452027
Sequence 6254	AA398732	Se	quence 6306	AA452352
Sequence 6255	AA399275	Se	quence 6307	AA453352
Sequence 6256	AA400696	Se	quence 6308	AA453651
Sequence 6257	AA401206	Se	quence 6309	AA453719
Sequence 6258	AA402115	Se	quence 6310	AA453784
Sequence 6259	AA403120	Se	quence 6311	AA454187
Sequence 6260	AA403121	Se	quence 6312	AA454962
Sequence 6261	AA403188	Se	quence 6313	AA455668
Sequence 6262	AA404444	_	quence 6314	AA455894
Sequence 6263	AA416736	Se	quence 6315	AA456390
Sequence 6264	AA417876	Se	quence 6316	AA458761
Sequence 6265	AA418628	Se	quence 6317	AA458770
Sequence 6266	AA419263	Se	quence 6318	AA459167
Sequence 6267	AA419554	_	quence 6319	AA459210
Sequence 6268	AA420571	Se	quence 6320	AA461128
Sequence 6269	AA420633	Se	quence 6321	AA461467
Sequence 6270	AA420705	Se	quence 6322	AA461605
Sequence 6271	AA421682	Se	quence 6323	AA464133
Sequence 6272	AA421850	Se	quence 6324	AA464141
Sequence 6273	AA422060	Se	quence 6325	AA467750
Sequence 6274	AA423801	Se	quence 6326	AA468167
Sequence 6275	AA425407	Se	quence 6327	AA468404
Sequence 6276	AA425526	Se	quence 6328	AA468624
Sequence 6277	AA426578	Se	quence 6329	AA468839
Sequence 6278	AA427866	Se	quence 6330	AA469129
Sequence 6279	AA428008	Se	quence 6331	AA469303
Sequence 6280	AA428084	Se	quence 6332	AA469345
Sequence 6281	AA429014	Se	quence 6333	AA469406
Sequence 6282	AA429293	Se	quence 6334	AA476524
Sequence 6283	AA429493	Se	quence 6335	AA476758
Sequence 6284	AA430110	Se	quence 6336	AA479167
Sequence 6285	AA431677	Se	quence 6337	AA479280
Sequence 6286	AA432103	Se	quence 6338	AA479287
Sequence 6287	AA433869	Se	quence 6339	AA479646
Sequence 6288	AA433988	Se	quence 6340	AA481078
Sequence 6289	AA435565	Se	quence 6341	AA481316
Sequence 6290	AA437224	Se	quence 6342	AA482539
Sequence 6291	AA442184	Se	quence 6343	AA482724

Page 61 of 74

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Sequence 6344	AA483214	Sequence 6396	443201U4
Sequence 6345	AA485928	Sequence 6397	4A528123
Sequence 6346	AA487448	Sequence 6398	AA528202
Sequence 6347	AA487747	Sequence 6399	AA531221
Sequence 6348	AA488633	Sequence 6400	AA531312
Sequence 6349	AA492035	Sequence 6401	AA531509
Sequence 6350	AA492280	Sequence 6402	AA531563
Sequence 6351	AA493222	Sequence 6403	AA531602
Sequence 6352	AA493331	Sequence 6404	AA533057
Sequence 6353	AA493448	Sequence 6405	AA533648
Sequence 6354	AA493512	Sequence 6406	AA534235
Sequence 6355	AA493956	Sequence 6407	AA534281
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Sequence 6357	AA494493	Sequence 6409	AA535377
Sequence 6358	AA495759	Sequence 6410	AA536036
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Sequence 6363	AA502979	Sequence 6415	AA552715
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Sequence 6365	AA504334	Sequence 6417	AA554735
Sequence 6366	AA504953	Sequence 6418	AA557888
Sequence 6367	AA505523	Sequence 6419	AA559906
Sequence 6368	AA505953	Sequence 6420	AA564484
Sequence 6369	AA507442	Sequence 6421	AA564505
Sequence 6370	AA507777	Sequence 6422	AA573742
Sequence 6371	AA508889	Sequence 6423	AA573762
Sequence 6372	AA512974	Sequence 6424	AA575882
Sequence 6373	AA514938	Sequence 6425	AA576425
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Sequence 6379	AA523252	Sequence 6431	AA578976
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Sequence 6383	AA523902	Sequence 6435	AA583567
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Sequence 6385	AA524778	Sequence 6437	AA587140
Sequence 6386	AA525419	Sequence 6438	AA587226
Sequence 6387	AA526515	Sequence 6439	AA587630
Sequence 6388	AA526591	Sequence 6440	AA592908
Sequence 6389	AA526886	Sequence 6441	AA594681
Sequence 6390	AA527187	Sequence 6442	AA594682
Sequence 6391	AA527272	Sequence 6443	AA595471
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Sequence 6393		Sequence 6445	AA602957
Sequence 6394	AA527805	Sequence 6446	AA603041
Sequence 6395		Sequence 6447	AA603135
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Page 62 of 74

Table 4-1

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Sequence 6448	AA604474	Sequence 6500	AA699999
Sequence 6449	AA604497	Sequence 6501	AA702398
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Sequence 6451	AA609094	Sequence 6503	AA705447
Sequence 6452	AA609436	Sequence 6504	AA705736
Sequence 6453	AA610476	Sequence 6505	AA705743
Sequence 6454	AA613916	Sequence 6506	AA705813
Sequence 6455	AA613933	Sequence 6507	AA706241
Sequence 6456	AA614105	Sequence 6508	AA707490
Sequence 6457	AA614525	Sequence 6509	AA708143
Sequence 6458	AA618484	Sequence 6510	AA708195
Sequence 6459	AA620750	Sequence 6511	AA713687
Sequence 6460	AA620995	Sequence 6512	AA714934
Sequence 6461	AA622344	Sequence 6513	AA716751
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Sequence 6463	AA625682	Sequence 6515	AA723679
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Sequence 6465	AA628526	Sequence 6517	AA724868
Sequence 6466	AA629534	Sequence 6518	AA729119
Sequence 6467	AA629608	Sequence 6519	AA732326
Sequence 6468	AA631868	Sequence 6520	AA740261
Sequence 6469	AA631903	Sequence 6521	AA741056
Sequence 6470	AA633794	Sequence 6522	AA742231
Sequence 6471	AA634039	Sequence 6523	AA744367
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Sequence 6473	AA634379	Sequence 6525	AA764981
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Sequence 6475	AA639317	Sequence 6527	AA772603
Sequence 6476	AA640105	Sequence 6528	AA773009
Sequence 6477	AA640141	Sequence 6529	AA773362
Sequence 6478	AA641623	Sequence 6530	AA773374
Sequence 6479	AA642471	Sequence 6531	AA775914
Sequence 6480	AA648508	Sequence 6532	AA778038
Sequence 6481	AA649193	Sequence 6533	AA778109
Sequence 6482	AA652846	Sequence 6534	AA778596
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Sequence 6484	AA654557	Sequence 6536	AA810159
Sequence 6485	AA654675	Sequence 6537	AA812193
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Sequence 6489	AA662702	Sequence 6541	AA826649
Sequence 6490	AA662822	Sequence 6542	AA827331
Sequence 6491	AA664480	Sequence 6543	AA831795
Sequence 6492	AA668224	Sequence 6544	AA831876
Sequence 6493	AA668692	Sequence 6545	AA835377
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Sequence 6495	AA670280	Sequence 6547	AA836488
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Sequence 6497	AA677282	Sequence 6549	AA845409
Sequence 6498	AA680052	Sequence 6550	AA845442
Sequence 6499	AA682910	Sequence 6551	AA846152
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Page 63 of 74

Sequence 6552	AA846576		A1028507
Sequence 6553	AA854071	Sequence 6605	A1028516
Sequence 6554	AA855093	Sequence 6606	A1033912
Sequence 6555	AA862250	Sequence 6607	A1038038
Sequence 6556	AA876668	Sequence 6608	A1038986
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Sequence 6558	AA884972	Sequence 6610	A1039636
Sequence 6559	AA884992	Sequence 6611	A1040598
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Sequence 6561	AA886957	Sequence 6613	A1042152
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Sequence 6569	AA907003	Sequence 6621	A1054023
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Sequence 6572	AA921906	Sequence 6624	AI064691
Sequence 6573	AA923172	Sequence 6625	AI074143
Sequence 6574	AA926926	Sequence 6626	AJ074389
Sequence 6575	AA927818	Sequence 6627	AI075893
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Sequence 6578	AA935181	Sequence 6630	Al082272
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Sequence 6602	AI018773	Sequence 6654.	Al142095
Sequence 6603	AI022058	Sequence 6,655	Al142558

Page 64 of 74

Sequence 6656		Sequence 6708	AJ267849
Sequence 6657		Sequence 6709	Al269060
Sequence 6658	Al148251	Sequence 6710	Al272941
Sequence 6659	Al148561	Sequence 6711	
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Sequence 6671	Al198986	Sequence 6722	Al302102
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Sequence 6673	Al199681	Sequence 6724	AI306457
Sequence 6674	Al200798	Sequence 6725	Al307419
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Sequence 6683	Al207618	Sequence 6734	Al342409
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Sequence 6685	AI214272	Sequence 6736	Al342669
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Sequence 6690	Al221894	Sequence 6741	Al352141
Sequence 6691	Al222329	Sequence 6742	Al357472
Sequence 6692	Al222789	Sequence 6743	AI358685
Sequence 6693	Al222806	Sequence 6744	AI360710
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Sequence 6697	Al253319	Sequence 6748	Al376840
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Sequence 6706	A1267652	, Sequence 6757	Al420898
Sequence 6707	A1267656	Sequence 6758	Al423180
-cdacine 0101	Al267664		Al432306

Page 65 of 74

Sequence 6760 Al432462	Sequence 6812 Al681405
Sequence 6761 Al433976	Sequence 6813 Al683298
Sequence 6762 Al435153	Sequence 6814 Al684170
Sequence 6763 Al435945	Sequence 6815 Al686139
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Sequence 6770 Al475476	Sequence 6822 AI701140
Sequence 6771 Al523940	Sequence 6823 Al703256
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	Sequence 6825 Al709186
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Ocqueries et :	Sequence 6827 AI718825
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	Sequence 6830 AI735314
004_00.00	Sequence 6831 Al735683
	Sequence 6832 AI739102
	Sequence 6833 AI741354
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Ocquerios er en	Sequence 6835 Al742468
	Sequence 6836 AI742912
Sequence 6784 Al567102	Sequence 6837 Al743852
Sequence 6785 Al569556	Sequence 6838 AI744509
Sequence 6786 Al569819	Sequence 6839 AI749547
Sequence 6787 Al569955	Sequence 6840 AI751364
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Sequence 6810 Al672201	Sequence 6863 AI82755
Sequence 6811 Al672868	26dnsucs good - Vigs 1 ag

Page 66 of 74

Table 4-1

Sequence 6864	Al828670	Sequence 6916	AL121547
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Sequence 6866	Ai861878	Sequence 6918	AW003692
Sequence 6867	Al879179	Sequence 6919	AW005103
Sequence 6868	Al879995	Sequence 6920	AW008213
Sequence 6869	Al887632	Sequence 6921	AW015683
Sequence 6870	Al888438	Sequence 6922	AW020479
Sequence 6871	A1890240	Sequence 6923	AW022300
Sequence 6872	Al910552	Sequence 6924	AW022756
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Sequence 6874	AI914128	Sequence 6926	AW051386
Sequence 6875	Al914133	Sequence 6927	AW052140
Sequence 6876	Al921434	Sequence 6928	AW062933
Sequence 6877	Al922855	Sequence 6929	AW067913
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Sequence 6879	Al923978	Sequence 6931	AW071903
Sequence 6880	Al929509	Sequence 6932	AW080160
Sequence 6881	AI937296	Sequence 6933	AW083804
Sequence 6882	A!940170	Sequence 6934	AW090544
Sequence 6883	AI948503	Sequence 6935	AW104561
Sequence 6884	Al954458	Sequence 6936	AW135430
Sequence 6885	Al961840	Sequence 6937	AW138413
Sequence 6886	Al963623	Sequence 6938	AW151295
Sequence 6887	A1970594	Sequence 6939	AW157047
Sequence 6888	Al972279	Sequence 6940	AW157303
Sequence 6889	Al973109	Sequence 6941	AW160731
Sequence 6890	A1983220	Sequence 6942	AW161491
Sequence 6891	AL035731	Sequence 6943	AW168960
Sequence 6892	AL035985	Sequence 6944	AW169876
Sequence 6893	AL036123	Sequence 6945	AW170485
Sequence 6894	AL036169	Sequence 6946	AW179006
Sequence 6895	AL036483	Sequence 6947	AW183759
Sequence 6896	AL036801	Sequence 6948	C06419
Sequence 6897	AL036988	Sequence 6949	D55192
Sequence 6898	AL037267	Sequence 6950	D58694
Sequence 6899	AL037471	Sequence 6951	F05508
Sequence 6900	AL037800	Sequence 6952	F06621
Sequence 6901	AL039253	Sequence 6953	F06739
Sequence 6902	AL040873	Sequence 6954	F12493
Sequence 6903	AL041050	Sequence 6955	F22763
Sequence 6904	AL041780	Sequence 6956	H00112
Sequence 6905	AL042316	Sequence 6957	H03785
Sequence 6906	AL044447	Sequence 6958	H12527
Sequence 6907	AL044649	Sequence 6959	H13339
Sequence 6908	AL045213	Sequence 6960	H16455
Sequence 6909	AL048699	Sequence 6961	H20086
Sequence 6910	AL049012	Sequence 6962	H20888
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Sequence 6912	AL118621	Sequence 6964	H23837
Sequence 6913	AL119462	Sequence 6965	H39906
Sequence 6914	AL120798	Sequence 6966	H39960
Sequence 6915	AL120840	Sequence 6967	H45550
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Page 67 of 74

Sequence 6968 H50566	Sequence 7020 R89611	
Sequence 6969 H57382	Sequence 7021 R93759	)
Sequence 6970 H59731	Sequence 7022 R98423	į.
Sequence 6971 H59791	Sequence 7023 T05006	
Sequence 6972 H70670	Sequence 7024 T05635	i
Sequence 6973 H75695	Sequence 7025 T10307	
Sequence 6974 H78954	Sequence 7026 T27258	\$
Sequence 6975 H81659	Sequence 7027 T28083	3
Sequence 6976 H83574	Sequence 7028 T31772	2
Sequence 6977 H84729	Sequence 7029 T6142	i .
Sequence 6978 H86010	Sequence 7030 T65998	₿.
	Sequence 7031 T7237	
Sequence 6979 L49115 Sequence 6980 M85423	Sequence 7032 T7894	В
Sequence 6981 N25145	Sequence 7033 T8370	9
Sequence 6982 N26359	Sequence 7034 T8494	3
Sequence 6983 N26421	Sequence 7035 U6955	<b>:9</b>
ocquee	Sequence 7036 W005	56
	Sequence 7037 W031	70
004201100 0000	Sequence 7038 W034	71
Ocquories sees	Sequence 7039 W054	72
Sequence 6987 N40852 Sequence 6988 \ N41386	Sequence 7040 W071	09
	Sequence 7041 W198	99
Ocquones sees	Sequence 7042 W252	22
	Sequence 7043 W259	70
Sequence 6991 N44001	Sequence 7044 W274	140
Sequence 6992 N45940	Sequence 7045 W285	596
Sequence 6993 N52271	Sequence 7046 W28	729
Sequence 6994 N52554 Sequence 6995 N63991	Sequence 7047 W28	994
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00400	Sequence 7049 W40	294
20420	Sequence 7050 W45	179
004201100	Sequence 7051 W45	265
00400	Sequence 7052 W51	959
00400	Sequence 7053 W56	523
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Ocque.iee		3793
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	Sequence 7062 W9	9251
		9471
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Sequence 7016 R70995		3002386
Sequence 7017 R71431		3002387
Sequence 7018 R74597		3002451
Sequence 7019 R80852	Ocqueilos (O/)	

Page 68 of 74

Sequence 7072		Sequence 7124	<b>AE</b> 000000
Sequence 7073		Sequence 7125	
Sequence 7074		Sequence 7126	
Sequence 7075		Sequence 7127	AF007142
Sequence 7076		Sequence 7127	AF007791
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Sequence 7078	AB011079	Sequence 7129	AF013717
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Sequence 7084		Sequence 7135	AF020202
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Sequence 7087	AB016068	Sequence 7138	AF021819
Sequence 7088	AB017004	Sequence 7139	AF022215
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Sequence 7098	AB020686	Sequence 7149	AF038452
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Sequence 7100	AB020692	Sequence 7151	AF039692
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Sequence 7103	AB021663	Sequence 7154	AF044321
Sequence 7104	AB023145	Sequence 7155	AF044670
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Sequence 7105 Sequence 7106	AB023173	Sequence 7157	AF044956
	AB023205	Sequence 7158	AF045167
Sequence 7107	AB023210	Sequence 7159	AF045229
Sequence 7108	AB023230	Sequence 7180	AF047020
Sequence 7109	AB026190	Sequence 7161	AF049910
Sequence 7110	AB027013	Sequence 7162	AF051323
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Sequence 7114	AB029004	_	AF054187
Sequence 7115	AB032983	_	AF054838
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Sequence 7122	AD001528		AF062323
Sequence 7123	AF000381		AF065388
		oequence /1/5 /	AF067171

Page 69 of 74

Sequence 7176 AF068235	Sequence 7228 AF131758
Sequence 7177 AF070523	Sequence 7229 AF131763
Sequence 7178 AF070539	Sequence 7230 AF131814
Sequence 7179 AF070555	Sequence 7231 AF131838
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Sequence 7181 AF070561	Sequence 7233 AF132000
Sequence 7182 AF070650	Sequence 7234 AF141968
Sequence 7183 AF070655	Sequence 7235 AF147330
	Sequence 7236 AF147331
Ocque: 100 T T T	Sequence 7237 AF147339
Sequence 1100	Sequence 7238 AF147395
Sequence 7186 AF075587 AF077030	Sequence 7239 AF151103
004-0	Sequence 7240 AF151109
Ocquorico	Sequence 7241 AF151803
Sequence : re-	Sequence 7242 AF151818
Ocquerios 1123	Sequence 7243 AF151840
Ouque.	Sequence 7244 AF151867
00420	Sequence 7245 AF151871
004201104	Sequence 7246 AF151878
Ouque	Sequence 7247 AF153608
Ocquerios	Sequence 7248 AF155095
Sequence : 100	Sequence 7249 AF155110
Sequence First	Sequence 7250 AF156965
Sequence : 100	Sequence 7251 AF159056
Sequerios , res	Sequence 7252 AF168956
Sequence (200	Sequence 7253 AF169797
Sequence rest	Sequence 7254 AF170583
00400.00	Sequence 7255 AF174605
Ocquerioe : =00	Sequence 7256 AF 176574
ocque	Sequence 7257 AF179274
Ocqueiloo 1220	Sequence 7258 AF182289
	Sequence 7259 AF182645
Ocquoiles . Les	Sequence 7260 AF188745
Ocqueino : ===	Sequence 7261 AF188746
Sequence 7209 AF086517 Sequence 7210 AF086557	Sequence 7262 AF195120
Sequence 7211 AF087999	Sequence 7263 AF201077
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Sequence 7225 AF 126782	Sequence 7278 AL049969
Sequence 7220 AF 120702 Sequence 7227 AF129927	Sequence 7279 AL050159
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Page 70 of 74

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Sequence 7281		Sequence 7332	
Sequence 7282		Sequence 7333	
Sequence 7283	AL050265	Sequence 7334	
Sequence 7284		Sequence 7335	
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Sequence 7294	AL110185	Sequence 7345	D87666
Sequence 7295	AL117423	Sequence 7346	D87735
Sequence 7296	AL117429	Sequence 7347	D87742
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Sequence 7298	AL117458	Sequence 7349	D89053
Sequence 7299	AL117621	Sequence 7350	D90209
Sequence 7300	AL117644	Sequence 7351	D90373
Sequence 7301	AL117666	Sequence 7352	E01650
Sequence 7302	AL122088	Sequence 7353 Sequence 7354	E02628
Sequence 7303	AL133074		E02822
Sequence 7304	AL133076	Sequence 7355 Sequence 7356	E06721
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Sequence 7313	D14812	Sequence 7365	J03799
Sequence 7314	D16911	Sequence 7366	J04164
Sequence 7315	D17039	Sequence 7367	J04205
Sequence 7316	D17554	Sequence 7368	J04208
Sequence 7317	D21262	Sequence 7369	J04443
Sequence 7318 Sequence 7319	D21853	Sequence 7370	J04794
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Sequence 7321	D26598	Sequence 7372	K03515
Sequence 7322	D28476	Sequence 7373	L00635
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Sequence 7325	D30655	Sequence 7376	L05093
Sequence 7326	D31883	Sequence 7377	L05779
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Sequence 7330	D42047	Sequence 7381	L10678
Sequence 7331	D43950	Sequence 7382	L16558
	- 10000	Sequence 7383	L19597

Page 71 of 74

	Sequence 7436 M74509	
Sequence 7384 L19605	Sequence 7436 M74509 Sequence 7437 M74777	
Sequence 7385 L20773	Sequence 7438 M76729	
Sequence 7386 L20941	Sequence 7439 M77830	
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Sequence 7388 L33404	Sequence 7441 M81757	
Sequence 7389 L34840	Sequence 7442 M83822	
Sequence 7390 L38486	Sequence 7443 M90054	
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Sequence 7392 L47276	Sequence 7445 M95724	
Sequence 7393 L49504	Sequence 7446 M97347	
Sequence 7394 L77964	Sequence 7447 S66196	
Sequence 7395 M10119	Sequence 7448 S66431	
Sequence 7396 M10941	Sequence 7449 S67815	
Sequence 7397 M11058	Sequence 7450 S69272	
Sequence 7398 M11353	Sequence 7451 S70154	
Sequence 7399 M12938	Sequence 7452 S70290	
Sequence 7400 M14200	Sequence 7453 S73498	
Sequence 7401 M14631	Sequence 7454 S73591	
Sequence 7402 M15178	Sequence 7455 S78203	
Sequence 7403 M16768	Sequence 7456 SB2081	
Sequence 7404 M16804	Sequence 7457 U03886	
Sequence 7405 M16942	Sequence 7458 U05875	
Sequence 7406 M17733	Sequence 7459 U08815	
Sequence 7407 M17885	Sequence 7460 U09559	)
Sequence 7408 M21154	Sequence 7461 U09820	)
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Sequence 7410 M21895	Sequence 7463 U14658	3
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Ocqueine	Sequence 7469 U1829	7
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	Sequence 7473 U2578	
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Sequence 7429 M55409	Sequence 7481 U372	
Sequence 7430 M58525	Sequence 7482 U376	
Sequence 7431 M61831	Sequence 7483 U388	
Sequence 7432 M63573	Sequence 7484 U39	_
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Sequence 7434 M69238	Sequence 7486 U40	
Sequence 7435 M74091	Sequence 7487 U47	142
Sequence : 100		

Page 72 of 74

Sequence 7488		Sequence 7540	X73902
Sequence 7489		Sequence 7541	X74801
Sequence 7490		Sequence 7542	X75593
Sequence 7491		Sequence 7543	
Sequence 7492	U65090	Sequence 7544	X75861
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	U96759	Sequence 7555	X85372
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Sequence 7514	X06617	Sequence 7566	Y13286
Sequence 7515	X07819	Sequence 7567	Z11241
Sequence 7516	X12597	Sequence 7568	Z14244
Sequence 7517	X12791	Sequence 7569	Z18538
Sequence 7518	X15187	Sequence 7570	Z26317
Sequence 7519	X15729	Sequence 7571	Z29330
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Sequence 7521	X52174	Sequence 7573	Z74615
Sequence 7522	X53605	Sequence 7574	
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Sequence 7525	X57352	Sequence 7577	N92483
Sequence 7526	X57766	Sequence 7578	Q13383
Sequence 7527	X57812	Sequence 7579	Q46089
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Sequence 7529	X59834		Q65676
Sequence 7530	X59841	Sequence 7581	Q75166
Sequence 7531	X63432	Sequence 7582	Q94780
Sequence 7532	X64330	Sequence 7583	T59274
Sequence 7533	X65644	Sequence 7584	V03517
Sequence 7534	X66276	Sequence 7585	V29653
Sequence 7535	X67698	Sequence 7586	V32447
Sequence 7536	X69117	Sequence 7587	V33197
Sequence 7537	X70326	Sequence 7588	V34263
Sequence 7538	X73114	Sequence 7589	V34271
Sequence 7539	X73608	Sequence 7590	V43612
-q-400 1000	A13000	Sequence 7591	V58360

Page 73 of 74

### Table 4-1

Sequence 7592 V	58628
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	<sup>7</sup> 58 <b>68</b> 4
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Sequence 7596 V	/62428
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	/87930
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	X26850
	X27262
	X37486
	X37532
	X39868
Sequence 7609	X40372
Sequence 7610	X51668
Sequence 7611	X52235
Sequence 7612	X59352
Sequence 7613	X91484
Sequence 7614	Z13088
Sequence 7615	Z15085
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Sequence 7617	Z16695
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Sequence 7619	727252
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Sequence 7623	AC29170
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Sequence 7627	AC32056
Sequence 7628	AC32097
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Sequence 7632	AC34190
Sequence 7633	AC003991
Sequence 7634	AC004130
Sequence 7635	
Sequence 7636	* *
Sequence 7637	
Sequence 7638	
Sequence 7639	
Sequence 7639	•
Sequence 7641	•
Sequence 7642	
Sequence 7643	3 AF064859
Sequence 7040	

Sequence 7644	AF103908
Sequence 7645	AL022097
Sequence 7646	AL031346
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Sequence 7648	AL079338
Sequence 7649	AP000527
Sequence 7650	S65371
Sequence 7651	U15177
Sequence 7652	Z83843
Sequence 7653	Z93020

Page 74 of 74

AAAAAAAAAA

Table 4-2

Sequence 1953:found in patent publication w099/64594
CCCTTAGCGTGGTCGCGGCCGAGGTACTCTTTCTCTCCCCTCCTCTGAATTTAATTCTTT
CAACTTGCAATTTGCAAGGATTACACATTTCACTGTGATGTATATTGTGTTGCAAAAAA
AAAAAGTGTCTTTGTTTAAAATTACTTGGTTTGTGAATCCATCTTGCTTTTTCCCCATTG
GAACTAGTCATTAACCCATCTCTGAACTGGTAGAAAAACATCTGAAGAGCTAGTCTATCA
GCATCTGACAGATGAAATTGGATGGTTCTCAGAACCATTTCACCCAGACAGCCTGTTTCT
ATCCTGTTTAATAAATTAATTTTTTGGGGTTCCTCTACCATGCCATAGCAAAACCCCCTGC
TTCCAAATTCTTGTCAACAATTAAAAAGTCTGNTGGACCTTGGAAGTTT

Sequence 1954:found in patent publication W099/40189
TTTTCCNTTTTTTTTTTTTTTTTTTTTTTTTCCCCNCCAAGCTTTATCTAGNCTTAGACT
TTTTAAAAAAGTTTGGGGGCAGATTCTGAATTGGCTAAAAGACATGCATTTTTAAAACTA
GCAACTCTTATTTCTTTCCTTTAAAAATACATAGCATTAAATCCCAAATCCTATTTAAAG

Page 1 of 8

Sequence 1956:found in patent publication W099/64594 CTCGAGCCGCCCCCCAGTGTGATGGGATATCTGCAGAATTCGCCCTTAGCCGTGGTCGC GGCCCGATGTA

Sequence 1958: found in patent publication W099/64594
CCCTTTCGAGCGGCCCCGGGCAGGTACTATATGGACACCAGCATGGAGCAGAAAATGA
GGGCAGCAGTGTCCAATCTCATTCCAGGTGAGAAGTTGCACAGTGTCCAATAGGTGCATA
CATCTCCTTAGTAAGTAGTTGTGATTAACAATGAAATAGAAATGAAAATATATTTTTT
ATTTATGTGTATTATATTTTTCAGGCAGCTAATAAGTTGGTAGGACATAATATTTTATTC
GTTGGGGACCTAATTATTTTATAAATTGAATGGTTAGATATTTCTTTTGGCCTAAGCCACC
ATGAAAAAATCACTGAGGCGCTAAGGGAAACATGAACTAAGAAGCCCTTCTGAGTCTCTG
TTTTCTCAACTGTAAGATGAAGAAACCTGCTCCACCTTCCTCTTGCGATTGTCGAACGGC
AGTTGAGATACTGCGCCAGAATGGACCCTTATTGATGGCCTAC

Sequence 1960:found in patent publication W099/64594 NGATATCTGCANAATTCGCCCTTTCGAGCGGCCGCCCGGGCAGGTACAACTTTTTAAATA GGGAATATGATAGCTTNGCATGGTGGTGTGCACCTATAGCCCCCACTGCCTGGAAAGCTG

Page 2 of 8

#### Table 4-2

GGGTGGAGAATCGCTTGAGTCCAGGAGTTTGAGGTTACAGTGATCCACGATCNNGCCAC TACACTCCAGCCTGGGCANCAGAGCAAGACCCTGTNTCAAAGCATAAAATGGAATNACAT ATCAAATGAAACANGGAAAATGAAGCTGA

CCCTTTCGAGCGGCCGCCCGGGCAGGTACTCAGACAAATACAACGTGCCCATTTCGTCAG
ATATTGCGCAGAACCAAGAATTTTATAAGAACGCAGAAGTTAGACCACCATTTACCGTAT
GCATNTTTAATTAAGGCAGGCCATTCTCGAATCTCCAGAAAAGCAGCTTAACACTAAATGA
GATCTATAACTGGTTCACACGAATGTTTGCTTACTTCCGACGCAACGCGGCCACGTGGAA
GAATGCAGTGCGTCATAATCTTAGTCTTCACAAGTGTTTTGTGGCGAGTAGAAAACGTTA
AAGGGGGCCAGTATGGGACAGTGGATTGAAGTANGAATTCCAAAAAACGAAGGGCCCAAAAA
GGATCAGTGGGTAACCCTTT

Page 3 of 8

GTGGGTGGGTGGCCACCTAGAAGACACATGGTNGGTTGTGCAACATAAAATCCCCTTCAAGCGGACTCAGTGG

Sequence 3729: found in patent application W099/64594 CCCTTAGCGTGGTCGCGGCCGAGGTACAGAATTTATTATGAAATAGCTTAATGGCAAGTG GTAATTTAGAAGAATTAACAGTTATCAGATAGGAGAATTTATTAAAATATTTAAAAATTTGATAAATTCTTGAAGCCCTTTTACACAAGTAATTTCTATAATTTGATTGTAATCAAAGTATAA TATACCTTGTTACTATTATCAGGATTAATTTTTGAAAGTAGAATTCCTTAATCAAGCCAA GGTTAATGCTGCTTTATAAGGAAATTAATCAGGTAGTTTAACACTAGAGCTCATTAGCCCA ACCTGTATGTAGCACAAAAATAAATCATTCTCTGATAAATNCCNTATTAAAATAATTTTTTTAATTCATACCTTTTTAAAATAA

Page 4 of 8

GAACCGGGGTTATTGGCCATAAATGGGNCCCCCAAATGGGTAANCCAAACTTTAAAAAAN ATGANCCGAAANCAAAGGGNAAAATCTGGGTGGCCCTTCCATTTNATTNTNGTANNNCAA **AGGGAACCTGGNNNAAGGGGGGCCTTTTNAAAAAA** Sequence 3732: found in patent application W099/64594 CCCAAAGCTTTATCTGTCTTGACTTTTTAAAAAAGTTTGGGGGCAGATTCTGAATTGGCT TTAAATCCCAAATCCTATTTAAAGACCTGACAGCTTGAGAAGGTCACTACTGCATTTATA GGACCTTCTGGTGGCTCTGTTACGTTTGAAGTCTGACAATCCTTGAGAATCTTTGCA TGCAGAGGAGGTAAGAGGTATTGGATTTTCACAGAGGAAGAACACAGCGCANAATGAAGG GCCAGGCTTACTGAGCTNTCCAGTGGAGGGCTCATGGGTGGGACATGGAAAANAAGGCAG CCTANGCCCTGGGGAGCCCANTCCACTGAGCAAGCAAGGGACTGANTGAGCCTTTTGCAG GAAAAGGCTTAAAAAAAGGAAAACCATTCTAAAACACAACAAGGAAACTGTCCAAATGCT TTGGGAACTGGGTTTATTTGCCTATTATGGGGTCCCCAAAAA Sequence 3733:found in patent application W098/39448 CCCTTAGCGTGGTCGCGGCCGAGGTACTACAAAAACAGAATAATTTTGAAGTTTTAGAAT AAATGTAATATTTACTATAATTCTAAATGTTTAAATGCTTTTCTAAAAATGCAAAACT ATGATGTTTAGTTGCTTTATTTTACCTCTATGTGATTATTTTTCTTAATTGTTATTTTTT ATAATCATTATTTTTCTGAACCATTCTTCTGGCCTCAGAAGTAGGACTGAATTCTACTAT TGCTAGGTGTGAGAAAGTGGTGGTGAGAACCTTAGAGCAGTGGAGATTTGCTACCTGGTC TGTGTTTTGAGAAGTGCCCCTTAGAAAGTTAAAAGAATGTAGAAAAGATACTCAGTCTTA ATCCTATGCAAAAAAAAATCAAGTAATTGTTTTCCTATGAGGAAAATAACCATGAGCTG TATCATGCTACTTAGCTTTTATGTAAATATTTCTTATGTCTCCTCTATTAAGAGTATTTA ATTTAACCCAATTGTCTCATATAAGGTTTTATTTGAATTG Sequence 4144: found in patent application W099/64594 CCCTTTCGGCGGCCCGGCCAGGTACCTGGGGTCTCAGGGTTGCTCTGGGCCTGATC ATCCACTCAGATCTGTAAGGAGGATTTGCAGGATCCATTTAGAAAGATCCTCCCTTACTT CCACAAGCATGGCCTTTGGCTCTTAAATACCTGTGCTGGGGTTTTGTAATTATAGAAACA CAGGGCTCAGGCCTAGAAAATCTAGTTTTCTGAAATCGCTAGCCAGCAACAGCACTGAG ATGGCCATCCCAGAAACAAGGCCAACACAGAAGCCACCATAAAGGCCGCTGGAGGTTGGG ACAAAGAGATCCTTGCTGTCCTTACAGACCCCCTGACTTCCAAGGAGCTCCCCTCTTACC CAGCCTGGCCTGCCTTCTCCACAGGGTAGCTGATCGTCAGCATCATCTTCAATGGTGTTG CCAAAAGCACTCAANTGCTCCTGCCATCCCTGTCCATCTTCAACATGAAAAGGAGAGGTT Sequence 4145: found in Patent application W099/64594 GGCTCCCCAAAGCTTTATCTGTCTTGACTTTTTAAAAAAGTTTGGGGGCANATTCTGAAT TAGCATTAAATCCCAAATCCTATTTAAAG Sequence 4146:found in Patent application W099/64594 ATCTGCAGAATTCGCCCTTANCGTGGTCGCGGCCGAGGTACTTTTT Sequence 4147: found Patent application W099/63088 CCCTTAGCGTGGTCGCGGCCCGAGGNACCATGATATCATGTATCCTGCTTGGACATTNTG GGAAAGGGGGACCAGCTGGTTGGCCAATTTATCCTACAGGTCTTGGACGGNGGGACCTCT TCAAAAGAAGATCTGGTAAGGGCAAGCAGCACAAGTGGCCATGGAAAAAAGAAAAACTCTA CANNATATTTCCGAGGATCAAGGACAAGTCCAAAAACGAGATCCTCTCATTCTTCTCTC GGAAAAACCCAAAACTTGTTTGATGCAGAATACACCAAAAACCAGCCTGGAAATCTATGA AAGATCCTTAGGAAANGCCAGCTGCTAAGGATGTCCATCTTGTGGATCACTGCAAATACA AGTATCTGTTTAATTTTCGAGGCGTAACTGCAAGTTTTCCGGTTAAACACCTCTTCCTGT GTGGCTCACTTGGTTTCCATGTTNGGTGATGAAGTGGCTAGAATTCTTCTATCCACAGCT TGAAGCCNTGGGTTCACTATATTCCAAGTCAAAACAGATCTCTTCAATGTCCAANAGCTG TTACAATTTTGAAAAACCAAATGATGATGTAACTTCAAGAAAATTGCTTGAAAGGGGGAA

Sequence 4148:found Patent application W099/64594 CCCTTAGCGTGGTCGCGGCCGAGGTACCACAAAAACAGAATAATTTTGAAGTTTTAGAAT

Page 5 of 8

#### Table 4-2

Sequence 7622: found in patent application W099/46374

Sequence 7623:found in patent application W099/51732
CGAGGTACGTTCTTCTCCAGTTTCTCTACCCTTAGCTTATCCTCACCCTCATTTTGC
CCTGCTGGCTGCTCAAACTATGCAACAGATTCGGCATCCTCGCTTACCCATGGCCCAGTT
TGGAGGAACCTTCTCACCTTCTCCTAAACACATGGGGACCATTCCCAGTGAGACCTGTGAA
TCCTGGCAACACAAATAGCTCTCCAAAGCATAATAACACAAGCCGTCTACCTAACCAGAA
CGGGACTGTTTTACCCTCAGAGTCTGCTGGACTAGCTACTGCCAGTTGTCCTATCACTGT
CTCTTCTGTAGTTGCTGCCAGTCAGCAACTGTGTCTCACTAATACCCGGACTCCTTCATC
AGTCAGAAAGCAGTTGTTTGCCTGTGTGCCTAAGACAAGTCCTCCAGCAACAGTGATTTC
TTCTGTGACAAGCACTTGTAGTTCCCTGCCTTCTGTCTCCTCTGCACCTATCACTAGCGG
GCAAGCTCCCACCACATTTCTACCTGCAAGTACCTGCCCG

Sequence 7624:found in patent application W099/42470 CCGCGGTGGCCGCCCGGGCAGGTACCGATACAGAAGAAGCTGCGTTTTGAAGACACC CTGGAGTTTGTAGGCTTGATGCGAAGATGGCTGAGGAATCCTCCTCCTCCTCA TCTTCACCAACTGCTGCAACATCTCAGCAGCAGCAACTTAAAAATAAGAGTATATTAATC TCTTCTGTGGCTTCGGTGCATCATGCAAACGGCCTAGCCAAATCTTCTACCACCGTCTCT AGCTTTGCTAACAGCAAACCTGGCTCTGCTAAGAAGTTAGTGATCAAGAACTTTAAAGAT AAGCCTAAATTACCAGAAAAACTACACAGATGAAACCTGGCAAAAACTGAAAGAAGCAGTG GAAGCTATTCANAATAGTACCT

Sequence 7626: found in patent application W099/57132 CGCCCGGGGCAGGTACCTTAGTGAGGCTCAAAAGGATTCTTTTGGGTCTATTTTACGCCT TATCTTTGAAATTCACCACTCGGGTGAGAAAGGTGACATTGTAGTCTTTCTGGCCTGTGA ACAAGATATTGAGAAAGTCTGTGAAACTGTCTATCAAGGATCTAACCCAGATCT TGGAGAACTGGTGGTTGTTCCTTTGTATCCAAAAGAGAAATGTTCATTGTTCAAGCCACT CGATGAAACAGAAAAAAAAGATGCCAAGTTTATCAAAGAAGAGTGGTGTTAACTACTACTC TGGAGAGTTTTTGATCTGGAGCAACTCAGTCAGATTTGTTATCGATGTGGGTGTGAAAA GAAAAAAAGGTGTCCTCGGC

Page 7 of 8

#### Table 4-2

CAGAAATGGCTGTTTCTGAGCTACCTGGTAACCCCAACGCTGTCTGGACAGTGCGTCGAC ACATTGAAGATGAGTTTGATGCCTACATCATTGTGTCTTTCGTGAATGCCACCCTAGTGT TGGCCATTGGAGAAACTGTAGAAGAAGTGACTGACTCTNGGTTCCTGGGGACCAC Sequence 7629: found in patent application W099/47669 TCCGGANTATATGTTGTATTCGGCTGGGTCGAGGGTCTCAGGCAGAGTGCGCAGGCTCGA CGGCTTATACTTTGGGAACGACATCTTGGCGAACCAGGGCGCAATTGCGCCTGCGCGATT CTGAGGCCCTTTGTCTATGCTGACCTTCAGCTTCCCCCGCGTACCTGCCCG Sequence 7630: found in patent application W099/57144 CCGCGGTGGCGCCCCCGGNCAGGTACGCGGGGCTGGGCGCGGGGAACTGAAAGCCGGA AGGGGCAAGACGGGCTCAGTTCGTCATGGGGCTGTTTGGAAAGACCCAGGAGAAGCCGCC CAAAGAACTGGTCAATGAGTGGTCATTGAAGGTAAGAAAAGGAAATGAGAGTTGTTGACA Sequence 7631: found in patent application W099/53040 CGAATTGGAGCTCCACCGGGGTGGCGGCCCGGCCCGGGCAGGTACTTTTTGCTTTAGCAG TAAAAAGGAATGCTCTGGTTTTTTAACTGGCTCCTTGAGGAAGCCAGAAGATGGCATCTG CAGTGGAAGTAGGCACCACCCTCAGATCACGAGGCCCCTAGGAAGAAGCCTGTGGAAAT CAGCCTGTGATGTGTGGTGGGCCAGCACCTAGAANACACATGGTGGTTGTGCACATA **AATCCCTTCAGCGGACTCAGTGGAAGGATGAGAATTCTGAAAGTTCATGCACATTTTATT** GAGTAGGTAATATAAAAATGCTTTTTCTTTTTCATTTGTTACAAGTGCATGCTTTGANTG CCCACCATNTTCAGAAGTCAAAATTACAAAAGG

Sequence 7632: found in patent application W099/53040
CCGCGGTGGCGCCCCGGGCAGGTACGCGGGCGGGGATGGGAGCCACGCCTGAACTAG
AGTTCAGGCTGGATACATGTGCTCACCTGCTGCTCTTTGTCTTCCTAAGAGACAGAGAGTG
GGGCAGATGGAGGAGAAAAGTGAGGAATGAGTAGCATTCTGCCAAAAGGGCCC
CAGATTCTTAATTTAGCAAACTAAGAAGCCCCAATTAAAAAGCATTGTGGCTAAAGTCTAA
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GCAAGTTATCCAGGCTCGTAACCTGCATGCTGCCACCTTGAATCCCAAGGAGTATCTGCA
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Page 8 of 8

Table 5-1

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3	37408	R49695
4	37598	R51080
5	38554	R49731
6	40104	R54594
7	41850	R52786
8	42747	R61847
9	46365	H09774
10	46477	H09966
11	46694	H10192
<sup>'</sup> 12	47400	H10413
13	47652	H11433
14	50080	H17934
15	50486	H17618
16	50935	H18626
17	51378	H24011
18	51799	H23524
19	67237	T52700
20	69002	T54298
21	69935	T48692
22	81413	T60161
23	111812	T91225
24	120528	T95320
25	124447	R01094
26	127524	R08871
27	133864	R28660
28	136605	R35051
29	149288	R82644
30	151055	H02231
31	156962	R74321
32	196435	R91517
33	197651	R94504
34	200354	R96914
35	204483	H58234
36	205497	H57857
37	210697	
38	249070	H66840 H80063
39	258300	
40	258649	N30680
41	259950	N32226
42	266343	N32587
43	277173	N26556
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45	280375	N62936
46		N47113
70	293745	N65950

Page 1 (of 3 pages of Table 5-1)

## Table 5-1

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48	344126	W73753
49	344190	W69741
50	344721	W74673
51	346308	W74133
52	396350	AA758376
53	399101	AA733027
54	413148	AA707871
55	415473	W80447
56	415610	W80730
57	415619	W80741
58	417393	W88942
59	418150	W90164
60	418206	W90529
61	418279	W90323
62	428560	AA005401
63	431408	AA705839
64	432479	AA699494
65	449428	AA777883
66	451596	AA707080
67	451911	AA706969
68	455275	AA677575
69	490484	AA101632
70	501431	· AA115300
71	503741	AA131466
72	595318	AA164301
73	625875	AA188804
74	645565	AA204830
75	687297	AA235116
76	712950	AA282273
7 <b>7</b>	729953	AA412049
78	731218	AA416724
79	742049	AA401457
80	742569	AA400074
81	753745	AA410491
82	755304	AA436327
83	767289	AA418418
84	767823	AA418728
85	768217	AA424905
86	782233	AA431721
87	782503	AA431773
88	785663	AA449085
89	786525	AA452113
90	786602	AA478470
91	788180	AA453310
92	795499	AA454215
93	810873	AA459197
94	810987	AA485357
95	815183	AA481152

Page 2 (of 3 pages of Table 5-1)

Table 5-1

96	815556	AA456821
97	824723	AA488986
98	839081	AA487608
99	840878	AA482324
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101	845345	AA773478
102	884606	AA630006
103	897924	AA598808
104	950470	AA599102
105	1030855	AA621761
106	1049033	AA778675
107	1049185	AA620697
108	1049230	AA620715
109	1292432	AA718910
110	1323448	AA873604
111	1412481	AA845156
112	1416782	AA894557
113	1420370	AA857035
114	1468220	AA884897
115	1471841	AA873355
116	1472753	AA872402

Page 3 (of 3 pages of Table 5-1)

Table 5-2

Order	Clone	Gen Bank Accession Number
1	62 <del>9</del> 896	AA219045
2	897567	AA489611
3	856650	AA669314
4	897781	AA598517
5	882510	AA676460
6	586685	AA130579
7	769890	AA430382
8	525926	AA074511
9	714414	AA293215
10	812251	AA455056
11	192569	H41489
12	814989	AA465723
13	781089	AA430032
14	757265	AA426113
	359661	AA011096
15	768562	AA429297
16	811062	AA485441
17	878676	AA775355
18	740941	AA478298
19	684655	AA251770
20	842863	AA489261
21	878798	AA670408
22	281010	N47691
23	135221	R32952
24	491751	AA150500
25 25	770879	AA434403
26 27	290841	N71982
28	291341	N72263
29	281003	N50880
30	785368	AA476576
30 31	45582	H08120
32	739511	AA478066
33	770043	AA427570
33	624627	AA187351
35	416833	W86653
36	359982	AA063521
	669375	AA253464
37 38	744047	AA629262
39	451907	AA706968
	757244	AA426025
40 41	453107	AA700904
· ·	755578	AA419177
42	502518	AA156802
43	773286	AA425299
44	843321	AA489569
45	278687	N62924
46		· · · ·
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Page 1

Table 5-2

47	853006	AA668230 '
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49	757165	AA443950
50	230971	R96155
51	796624	AA460530
52	38740	R51273
53	868757	AA775325
54	825372	AA504507
55	746245	AA417761
56	. 292399	N68399
57	626716	AA191548
58	489106	AA056538
59	22074	T66180
60	85800	T72067
61	705064	AA279990
62	277740	N49589
63	35300	R43798
64	788256	AA454098
65	746007	AA482007
<b>6</b> 6	455263	AA677572
67	277112	N39611
68	259591	N32768
69	22773	R38613
70	2699 <b>97</b>	N24910
71	451918	AA706964
72	345626	W72051
73	51041	H18633
74	32299	R42685
75 	812266	AA455062
76 	41391	R56123
77	878253	AA775791
78 70	83920	T64312
79	740604	AA479795
80	770066	AA430545
81	196189	R92281
82	46367	H09959
83	7847 <b>72</b>	AA478542
84 85	246430	N53031
	310493	W31074
86 87	753162	AA400457
87 88	459941	AA779380
	200814	R98851
89	415102	W95001
90 91	489677	AA099568
	193892	H51765
92	359781	AA011320
93	812266	AA455062
94	277414	N34436
95	49284	H15533

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Table 5-2

	66317	T66816
96	239611	H79534
97		AA129777
98	502151	AA425450
99	773330	N24113
100	266819	R40970
101	29063	
102	785293	AA476543
103	824933	AA489040
104	345616	W72431
105	144816	R76281
106	291985	N73101
107	256907	N30096
108	487793	AA043501
109	840776	AA486082
110	199945	R97066
111	232946	H75599
112	197525	H52001
113	753467	AA406551
114	771303	AA443638
115	50879	H18424
116	341680	W60414
117	461727	AA682293
118	345793	W70128
119	486591	AA04299
120	345849	W70343
• =	452374	AA70087
121	241699	H91641
122		H99075
123	262060	1100070

Page 3 (of 3 pages of Table 5-2)

INTERNATIONAL SEARCH REPORT Int I Application No PCT/US 01/02318 A. CLASSIFICATION OF SUBJECT MATTER IPC 7 G01N33/574 C07K16/30 According to International Patent Classification (IPC) or to both national classification and IPC B. FIELDS SEARCHED Minimum documentation searched (classification system followed by classification symbols) IPC 7 G01N Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched Electronic data base consulted during the international search (name of data base and, where practical, search terms used) BIOSIS, CHEM ABS Data, MEDLINE, EPO-Internal, EMBL C. DOCUMENTS CONSIDERED TO BE RELEVANT Citation of document, with indication, where appropriate, of the relevant passages Category ° Relevant to claim No. Α WO 99 64594 A (CHIRON CORP) 16 December 1999 (1999-12-16) 1-55 abstract claims 1-15 WO 99 37811 A (AN GANG ; UROCOR INC (US); VELTRI ROBERT W (US)) Α 1-55 29 July 1999 (1999-07-29) abstract claims 1-32 -/--

X Further documents are listed in the continuation of box C.	X Patent family members are listed in annex.
<ul> <li>Special categories of cited documents:</li> </ul>	
"A" document defining the general state of the art which is not considered to be of particular relevance "E" earlier document but published on or after the international filling date "L" document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified) "O" document referring to an oral disclosure, use, exhibition or other means "P" document published prior to the international filing date but later than the priority date claimed Date of the actual completion of the international search	"T" later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention  "X" document of particular relevance; the claimed Invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone  "Y" document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other such documents, such combined with one or more other such documents, such combination being obvious to a person skilled in the art.  "&" document member of the same patent family
, and the medical section	Date of mailing of the international search report
19 July 2001	3 1. 10. 2001
larne and mailing address of the ISA European Patent Office, P.B. 5818 Patentlaan 2 NL - 2280 HV Rijswijk	Authorized officer
Tel. (+31-70) 340-2040, Tx. 31 651 epo nl, Fax: (+31-70) 540-3016	Gundlach, B

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	ation) DOCUMENTS CONSIDERED TO BE RELEVANT  Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Category °	Charlon of document, with indication, whose appropriate	
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ational application No. PCT/US 01/02318

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)  This International Search Report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:
This international Search Report has not been established in respect of certain claims under Article 17(0)(a) to the China
, and the following reasons:
1. X Claims Nos.:
because they relate to subject matter not required to be searched by this Authority, namely:
see FURTHER INFORMATION sheet PCT/ISA/210
2. Claims Nos.:
because they relate to parts of the International Assistant
an extent that no meaningful International Search can be carried out, specifically:
3. Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).
Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)
This International Searching Authority found multiple inventions in this international application, as follows:
see additional sheet
As all required additional access to
As all required additional search fees were timely paid by the applicant, this International Search Report covers all searchable claims.
As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
or any additional ree.
·
As only some of the required additional search fees were timely paid by the applicant, this International Search Report covers only those claims for which fees were paid, specifically claims Nos:
covers only those claims for which fees were paid, specifically claims Nos.:
No required additional search fees were timely paid by the applicant. Consequently, this International Search Report is
Colored by Claim's NOS.
see extra sheet invention 1.
The additional search fees were accompanied by the applicant's protest.
No protest accompanied the payment of additional search fees.
End of additional search fees.
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International Application No. PCT/US 01/02318

# FURTHER INFORMATION CONTINUED FROM PCT/ISA/ 210

This International Searching Authority found multiple (groups of) inventions in this international application, as follows:

1. Claims: Invention 1: Claims 1-55 (all in part)

Method of detecting/treating prostate cancer by determining the expression of a marker coded for by the sequence AA453310 (gene bank accession number).

2. Claims: Invention 2-19723: Claims 1-55 (all in part)

Method of detecting/treating prostate cancer by determining the expression of a marker coded for by one of the sequences listed in table 1-1 to 6.

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International Application No. PCT/US 01/02318

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